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Match Length
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-249-016-13408
US-09-249-016-13408
US-09-249-016-137497
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; TYPE: DNA
; ORGANISM: Human
US-09-949-016-80901
                                                                                                      CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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NUMBER OF SEQ ID NOS: 207012
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LOCATION: (1) ... (34408)

OTHER INFORMATION: n = A,T,C or G

US-09-949-016-14010
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                                                         SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 80901
LENGTH: 601
                                                                                                                                                                                                                                                                                    Sequence 80901, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: FOLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT FILING DATE: 1999-02-12
CURRENT FILING. DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
SEQ ID NO 10881
SEQ ID NOS: 28208
SEQ ID NO 10881
LENGTH: 366

TYPE- 1813-66

TYPE- 1813-66

THERAPEUTION OF SEQ ID NOS: 28208
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Best Local (
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ORGANISM: Candida albicans
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Pred. No. 1.
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Pred. No. 1e+02;
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CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 2809
SEQ ID NO 2809
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CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FASESEQ for Windows Version 4.0

TYPE: DNA

TYPE: DNA
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                                                                                                  US-09-248-796A-2809
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US-09-248-796A-2809/c
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                                     Matches
                                                                                                                                                                                                                                                                                         Sequence 2809, Application US/09248796A
PACENT NO. 6747137
GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
                                                                  Query Match
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 111397, A
Patent No. 6812339
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Best Local :
                                                                                                          TYPE: DNA ORGANISM: Candida albicans
                                                    Local
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2 ATTCCCCATCTCTCTTTCTTT 22
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19; Conserv
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                                 Conservative
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                                               Score 17.8;
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Pred. No. 1
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Pred. No. 1.2e+02;
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                                Mismatches
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                         1.2e+02;
2;
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                                                           Length 711;
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCH: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13408
LENGTH: 49378
RESULT 9
US-09-949-016-14083/c
; Sequence 14083, Application US/09949016
; Patent No. 6812339
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; Sequence 13408, Application US/09949016
; Patent No. 6812339
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; ORGANISM: Human
US-09-949-016-14816
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Best Local S
Matches 19
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LENGTH: 45086
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Patent No. 6812339
GENERAL INFORMATION:
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                       ORGANISM: Human
-09-949-016-13408
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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19; Conserv
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ilarity 90.5%;
Conservative
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Pred. No. 1.9e+02;
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LENGTH: 199471
; TYPE: DNA
; ORGANISM: Human
; PEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(199471)
; OTHER INFORMATION: n = A,T,C or
US-09-949-016-14083
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Sequence 69074, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 50/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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                                                                             Matches
                                                                                                                Query Match
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                                                                                                Best Local Similarity
                                                                                                                                                                           LENGTH: 601
TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: VENUER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
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331 Arrececarerererreer 313
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Similarity 90.5%;
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94.7%;
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Pred. No. 2.
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Pred. No. 1.8e+02;
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RESULT 11 US-09-949-016-137497/c US-09-949-016-137497, Application US/09949016 ; Patent No. 6812339 ; GENERAL INFORMATION:

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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT ETLING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILLING DATE: 2000-10-20
PRIOR PILLING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILLING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILLING DATE: 2000-09-08
PRIOR FILLING DATE: 2000-09-08
PRIOR FILLING DATE: 2000-09-08
SEQ ID NO 16528
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16528
LENGTH: 19503
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US-09-949-016-12500
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US-09-949-016-16528/c
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GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
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CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                   Sequence
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Best Local Similarity
                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                 Matches
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GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Human
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                                                                                                                             12500, Application US/09949016
o. 6812339
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94.7%;
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Pred. No. 1.8e+02;
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                                                                                                                                                                                                                                                                                                                                                          .6e+02
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                                         ASSOCIATED OF DETECTION
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                                       AND USES THEREOF
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-90-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2
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US-09-949-016-13769
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US-09-949-016-16092
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Sequence 13769, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,756
PRIOR PRICE STATE 2000-10-20
PRIOR PRICE STATE 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 16092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 681233
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18; Conservative
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Pred. No. 2.9e+02;
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Pred. No. 2.9e+02;
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PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR PILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13769
; LENGTH: 58844
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-13769

Query Match
Best Local Similarity 94.7%; Pred. No. 3e+02;
Matches 18; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 ATTCCCCATCTCTTTCT 20
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Db 16891 ATTCCCCATCTCTCTTCCT 16909

Search completed: February 28, 2005, 01:15:01
Job time: 7.47527 secs
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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         110987654321
                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                   Score
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                                                                                                                                                                                                                                                      Query
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1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
5: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
6: /cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/USOB_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/USOB_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/USOE_PUBCOMB.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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 11 US-09-909-317-1

US-10-027-632-190184

US-10-027-632-190184

US-10-027-632-190184

9 US-09-764-878-379

9 US-09-764-878-377

14 US-10-079-854-377

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17 US-10-180-375-7

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Sequence 1, Appli
Sequence 190184,
Sequence 190184,
Sequence 379, App
Sequence 377, App
Sequence 377, App
Sequence 377, Appli
Sequence 23, Appli
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9 1	17	8	7	17	3	19	17	13	1	17		18									13									18	17	15
US-10-203-319A US-09-938-842A-	US-10-203	US-10-767-701-	US-10-424-599-94947	US-10-027	US-10-027-632-10199	US-10-741	US-10-02	US-10-02	US-09-997-722-4		10-459-262F	719-993	10-240-485-	10-311-455-	10-027-632-	10-027-632-2069	-10-425-115-	-10-027-632-206	-10-027-632-206	-10-027-632	10-027-632-206	-10-674-124A-2346	10-424-599-3100	10-741	10-085	10-242-	10-741-600-176	-601	0-741-600-17	-10-433-793	-10-221-613-1	-10-311-455-14
Sequence 24, Appl Sequence 2730, Ap	e 22, App	12634	94947	Sequence 101998.	10199	26	e	e 1	43	<u>υ</u>		67	15	192	20691	æ	æ	O	æ	ø	Sequence 206913,	e 23460,	U	e 5616,	e 2643	e 26433,	e 17685,	e 5660	e 178	ю 	e 191	:-

ALIGNMENTS

RESULT 1 US-09-909-317-1

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; Sequence 1, Application US/09909317; Publication No. US20040152075A1

GENERAL INFORMATION:
APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
ITITLE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: 18810-82152
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR PELICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SOFTWARB: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
SEQ ID NO 1
LENGTH: 22
TYPE: DNA
ORGANISM: Homo sapiens
US-09-909-317-1
Query Match
Best Local Similarity 100.0%; Score 22; DB 11; Length 22;
Matches 22; Conservative 0; Mismatches 0; Indels 0; Gaps
Qy
1 GATTCCCCATCTCTCTTTTT 22
PRESULT 2
US-10-027-632-190184/c
US-10-027-632-190184/c
US-10-027-632-190184, Application US/10027632
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; TYPE: DNA
; ORGANISM: Human
US-10-027-632-190184
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; TYPE: DNA
; ORGANISM: Human
US-10-027-632-190184
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/16,002
PRIOR APPLICATION NUMBER: US 60/146,002
                                                                                                                                              NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID 190184
        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 190184, Application US/10027632
Publication No. US20030204075A9
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
                                                                                                                 LENGTH:
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 60/146,002
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  90.9%;
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100.0%; Pred. No.
Live 0; Mismatcl
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RESULT 6
US-09-764-878-377/c
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                                                                   Sequence 377, Application US/09764878 Patent No. US20020090615A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 19; Conserv
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SEQ ID NO 379
LENGTH: 32189
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NUMBER OF SEQ ID NOS: 428
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 379
LENGTH: 32189
TYPE: DNA
TYPE: DNA
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Patent No. US20020090615A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA121
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA121
CURRENT APPLICATION NUMBER: US/09/764,878
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Best Local :
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CURRENT FILING DATE: 2002-02-22
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 428
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA121C1
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CURRENT FILING DATE: 2001-01-17
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Similarity 100.0%; Pred. No.
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                                    and Antibodies
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RESULT 8
US-10-180-375-7/c
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, NAME/KEY: SITE
; LOCATION: (7464)
; OTHER INFORMATION: n equals a,t,g,
US-09-764-878-377
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Best Local S
Matches 19
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NUMBER OF SEQ ID NOS: 428
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 377
TENGTH: 32221
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                                                                                                                                                                                                                                 Sequence 7, Application US/10180375 Publication No. US20030126638A1
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SEQ ID NO 377
LENGTH: 32221
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Best Local Similarity
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA12IC1
CURRENT APPLICATION NUMBER: US/10/079,854
CURRENT FILING DATE: 2002-02-22
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ORGANISM: Homo sapiens
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NAME/KEY: misc feature
LOCATION: (7464)
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            ICANT: Shen, Bo
ICANT: Tarczynski, Mitchell C.
E OF INVENTION: Alteration Of Oil Traits In Plants
REFERENCE: BB1458 US NA1
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   APPLICATION NUMBER:
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19; Conserv
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                                                                                                                               Cahoon, Rebecca E. Famodu, Omolayo O. Harvell, Leslie T. Helentjaris, Timothy
                                                                                                 Li, Changjiang
Lowe, Keith
                                                                                Oliveira, Igor Cunha
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100.0%;
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100.0%; Pred. No. 1.6e+0
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US/10/180,375
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Pred. No. 1.6e+02;
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US-10-424-599-92893/c
Sequence 92893, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 222
SOFTWARE: Microsoft Office 97
SEQ ID NO 7
LENGTH: 1334
TYPE: DNA
ORGANISM: Vitis sp.
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APPLICANT: Allen,
APPLICANT: Cahoor
APPLICANT: Epelba
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APPLICANT: LA ROSA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223) B
CURRENT APPLICATION NUMBER: US/10/424,599
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TYPE: DNA
ORGANISM: Vitis
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TITLE OF INVENTION: Alteration Of Oil Traits
FILE REFERENCE: BB1458 US NA
CURRENT APPLICATION NUMBER: US/10/183,687
CURRENT FILING DATE: 2002-06-27
CURRENT FILING DATE: 2002-06-27
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APPLICANT:
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PRIOR FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 532
SOFTWARE: Microsoft Office 97
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Jones, Todd
Kinnon
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Oliveira, Igor Cunha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Allen, William
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90.9%; Pred. No. 1.7;
bive 0; Mismatches
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Pred. No. 1
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APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases As
TITLE OF INVENTION: Cytosine methylation
FILE REFERENCE: 5013 1014
CURRENT APPLICATION NUMBER: US/10/311,455
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: PCT/EP01/07537
PRIOR FILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: DC 10032529.7
PRIOR FILING DATE: 2000-06-30
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US-10-311-455-1419/c
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US-10-424-599-59036
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Chou Yihua
APPLICANT: Cap Yongwei
APPLICANT: Cap Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
FULE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 59036
LENGTH: 1309
                                                                                                                                                                                                    Sequence 1419, Application US/10311455
Publication No. US20030143606A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, Christian
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Best Local S
Matches 19
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Best Local
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Publication No. US20040031072A1
GENERAL INFORMATION:
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SEQ ID NO 92893
LENGTH: 284
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TYPE: DNA
ORGANISM: Glycine max
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LOCATION: (1)..(284)
OTHER INFORMATION: unsure at all n locations
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APPLICATION NUMBER: DE 10043826.1
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Pred. No. 2.5e+02;
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Pred. No. 2.4e+02;
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RESULT 14
US-10-433-793-11/c
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US-10-221-613-191/c
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Best Local .
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SEQ ID NO 191
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NUMBER OF SEQ ID NOS: 2424
SEQ ID NO 1419
LENGTH: 6351
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Best Local Similarity
Matches 19; Conserv
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CURRENT FILLING DATE: 2002-09-13
PRIOR APPLICATION NUMBER: PCT/EP01/02945
DE 1001387.00
DE 10019058.8
DE 10019058.8
DE 10032529.7
DE 10043826.1
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APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: OLEK, ALexander APPLICANT: PIEPENBROCK, C
                                                                                                                                                                                           FEATURE: OTHER INFORMATION: chemically treated genomic DNA
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TYPE: DNA
ORGANISM: Artificial Sequence
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o. US20040029123A1
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95.0%;
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Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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Sequence 11, Application US/10433793
Publication No. US20040142334A1
GENERAL INFORMATION:
APPLICANT: Epigenomics AG
TITLE OF INVENTION: Diagnose von mit Angiogenese

assoziierten Krankheiten

CURRENT APPLICATION NUMBER: US/10/433,793 CURRENT FILING DATE: 2003-06-06 NUMBER OF SEQ ID NOS: 212 SEQ ID NO 11

FILE REFERENCE

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RESULT 15
US-10-741-600-17886/c
US-10-741-600-17886/c
; Sequence 17886, Application US/10741600
; Publication No. US20050026169A1
; GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
FITLE OF INVENTION: MYCCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF
; TITLE OF INVENTION: MYCCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001499
; CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT ELING DATE: 2003-12-22
; NUMBER OF SEQ.ID NOS: 73997
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ.ID NO 17886
; LENGTH: 54016
Search completed: February 28, 2005, 06:52:25 Job time : 60.3236 secs
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                                                                                                                                                                                                                                                                                                 TYPE: DNA
CARANISM: Homo sapiens
FEATURE:
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(54016)
THER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables US-10-741-600-17886
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; TYPE: DNA
; TYPE: DNA
; ORANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-10-433-793-11
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Sequence 50, Appl Sequence 1279, Ap Sequence 157857, Sequence 157858, Sequence 157858, Sequence 157850,

Sequence Sequence Sequence

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Minimum
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Maximum Match 100%
Listing first 45 summaries
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

9: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

9: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-08-956-171E-38
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Sequence 186786, Appl Sequence 38, Appl Sequence 17554, A Sequence 11945, A Sequence 15909, A Sequence 15368, A Sequence 15368, A Sequence 15368, A Sequence 15368, A Sequence 1583, A Sequence 17999, A Sequence 17999, A Sequence 17993, A Sequence 17995, A Sequence 17997, A Sequence 50, Appl Sequence 
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Appli
Sequence 1722, Ap
Sequence 1722, Ap
Sequence 87851, A
Sequence 87852, A
Sequence 183202,
Sequence 183205,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Description
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US-08-956-171E-1722/c
^~~~ence 1722, Application US/08956171E
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SOPTWARE: FastSEQ for Wi
SEQ ID NO 2
LENGTH: 20
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-280-1818-2
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Best Local S
Matches 20
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: P07 41735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/280,1818
CURRENT FILING DATE: 1999-03-29
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CITY: Rockville
STATE: Maryland
COUNTRY: USA
ZIP: 20850
COMPUTER READABLE FORM:
                                                                                                                                                                                                  Gil H. Choi
Patrick S. Dillon
Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
                                                                                                                                                         NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                APPLICANT: Charles Kunsch
Gil H. Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AAATTGTGGTAATGACTGCA
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                                                                                                                                      ADDRESSEE: Human Genome
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US-09-667-878-50
US-09-265-480-50
US-09-834-759-50
US-09-590-751A-50
US-09-591-621-50
US-09-702-705-1279
US-09-713-25-1279
US-09-671-325-1279
US-09-680-824-1279
US-09-949-016-157859
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                                                                                                                                         Sciences,
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Length 20; Indels

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Gaps

CURRENT APPLICATION DATA:

SOFTWARE: ASCII

OPERATING SYSTEM: MSDOS version 6.2

MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33

APPLICATION NUMBER: US/08/956,171E

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RESULT 3
US-08-781-986A-1722/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1722, Application US/08781986A Patent No. 6737248
GENERAL INFORMATION:
   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                     NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PB248PP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (301) 309-843
INFORMATION FOR SEQ ID NO: 1722:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARDS. ACCTITETE TO STATE THE STATE OF THE STATE
                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                       TELEPHONE: (301)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Mark J. Hyman
REGISTRATION NUMBER: 46,789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
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TYPE: nucleic acid
STRANDEDNESS: double
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CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8: Human Genome Scier
9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Charles Kunsch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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1722:
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Pred. No. 4:
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GENERAL INFORMATION: Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION SEEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0

FERDING TO SEE TO S
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Patent No. 6812339

GENERAL INFORMATION:

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

PILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/331,498

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/311,498

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012
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; ORGANISM: Human
US-09-949-016-87851
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US-09-949-016-87851/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 87852, Application US/09949016
Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 87851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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ENGTH: 601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 325 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 84.0%; Local Similarity 90.0%; es 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          250 AAATTGAGGTAATGATTGCA 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57 AACGTGTGGTAATGACTGCA 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AAATTGTGGTAATGACTGCA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    h 84.0%;
Similarity 90.0%;
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Pred. No. 4
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Pred. No. 41;
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US-09-949-016-186785/c
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; Patent No. 6812339
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; ORGANISM: Human
US-09-949-016-183202
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                                                                                                                            ; ORGANISM: Human
US-09-949-016-186785
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLO01307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/247,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
Query Match
Best Local Similarity 90.0
Matches 18; Conservative
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GENERAL INFORMATION:
GENERAL INFORMATION:
J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 186785
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                                                                                                                                                                                                                                                                            WILLING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          156 AAATTGTGGTGATGATTGCA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            249 AAATTGAGGTAATGATTGCA 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18;
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Pred. No. 46;
0; Mismatches
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Pred. No. 46;
0; Mismatches
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Pred. No. 46
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OF DETECTION
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US-08-956-171E-38
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US-09-949-016-186786/c
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CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR PFLING DATE: 2000-10-20

PRIOR PFLING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR PILING DATE: 2000-09-08

PRIOR PR
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LENGTH: 601
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APPLICANT: VENTER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6593114
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence 186786,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                          ZIP: 20850

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 1.4

COMPUTER: HP Vectra 486/33

OPERATING SYSTEM: MSDOS version 6.2

SOFTWARE: ASCII Text

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/956,171E

PILING DATE: 20-Oct-1997

CLASSIFICATION: 4Unknown>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Charles Kunsch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            392 AAATTGTGGTGATGGCTGCA 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 Similarity 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AAATTGTGGTAATGACTGCA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 9410 Key West
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAATTGTGGTAATGACTGCA 20
                APPLICATION NUMBER: 60/009,861 FILING DATE: January 5, 1996 APPLICATION NUMBER: 08/781,986
                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84.0%;
nilarity 90.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patrick S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 . Choi
s. Dillon
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Pred. No. 46;
0; Mismatches
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                                                                                                                                                                                                                                                                                          3.50 inch, 1.4Mb storage
6/33
                                                                                                                                                                                                                                                                version 6.2
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Gaps

ATTORNEY/AGENT INFORMATION:

FILING DATE: January 3, 1997

NAME: Mark J. Hyman
REGISTRATION NUMBER: 46,789

PB248P1

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                                                                                        US-08-781-986A-38
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US-08-781-986A-38
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                                                                                                                                                   NAME: Benson, Bob
REGERENCE/DOCKET NUMBER: PB24/
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELESPHONE: (301) 309-8504
TELESPAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-956-171E-38
                              Matches
                                                           Query Match
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                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Diskette, 3.50 inch, 1.
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Charles Kun
TITLE OF INVENTION: St.
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                              STRANDEDNESS: dou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRIET: Marvin.
                                             Local
                                                                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U
ZIP: 20850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     y Match 84.0%;
Local Similarity 90.0%;
nes 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER:
                              18;
1 AAATTGTGGTAATGACTGCA 20
                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AAATTGTGGTAATGACTGCA 20
                                                                                                             nucleic acid
DEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AACGTGTGGTAATGACTGCA 22948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 23439 base pairs TYPE: nucleic acid
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                            Conservative
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VENTION: Staphylococcus aureus Polynucleotides and Sequences
                                          90.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                3.50 inch, 1.4Mb storage
                         ; Score 16.8; D; Pred. No. 87; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             314-1224
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Pred. No. 87;
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                                                   DB 4; Length 23439;
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                        Gaps
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; ORGANISM: Human US-09-949-016-11945
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US-09-949-016-11945/c
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                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11945
                                                                                                                                                                                                                                                                                                                                                                            Sequence 11945, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILLE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR PILING DATE: 2000-11-20
PRIOR PILING DATE: 2000-11-20
PRIOR PILING DATE: 2000-11-20
PRIOR PILING DATE: 2000-13-768
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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                                                                                            Query Match
Best Local :
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SEQ ID NO 1754
LENCTH: 62908
TYPE: DNA
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Patent No. 6812339
GENERAL INFORMATION:
                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                        LENGTH: 86439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: (1)...(62908)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: VENTER, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Human
                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity es 18; Conserv
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1 AAATTGTGGTAATGACTGCA 20
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                                                                                       Similarity
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                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                       84.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      84.0%;
                                                                                       Score 16.8; DB 4; Pred. No. 1.1e+02;
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Pred. No. 1e+02;
                                                                 Mismatches
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                                                                                     1.1e+02;
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                                                                                                     Length 86439;
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                                                                 Indels
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23617 AAATTGTGGTGATGATTGCA 23598

US-09-949-016-16990/c

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FORTIER FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
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                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ.ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16990
LENGTH: 86440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 16990, Application US/09949016 Patent No. 6812339
                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                                                 Matches
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-09-949-016-16990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
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PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                  NAME/KEY: misc_feature
LOCATION: (1)...(123463)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                ENGTH: 123463
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Local Similarity 90.0%;
nes 18; Conservative
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29756 AAATTGTGGTGATGGCTGCA 29737
                                                                                   18;
                                       μ
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                                     AAATTGTGGTAATGACTGCA 20
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                                                                                   Conservative
                                                                                                   84.0%;
                                                                               Score 16.8; DB 4;
Pred. No. 1.2e+02;
0; Mismatches 2;
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Pred. No. 1.1e+02;
0; Mismatches 2;
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                                                                                                                         Length 123463;
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Search completed: February 28, 2005, 01:15:05 Job time : 8.06842 secs
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US-09-949-016-12257/c
                                                                                                                                                                                                                                                         US-09-949-016-12257
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILLING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PELICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                     Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 12257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12257, Application US/09949016 Patent No. 6812339
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                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                    ORGANISM: Human
FEATURE:
NAMB/KEY: misc_feature
LOCATION: (1)...(129327)
OTHER INFORMATION: n = A,T,C
                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                         Local
                                                                                   68053 AAATTGTGGTGATGGCTGCA 68034
                                                                                                                                                                       18;
                                                                                                                             1 AAATTGTGGTAATGACTGCA 20
                                                                                                                                                                     h 84.0%;
Similarity 90.0%;
18; Conservative
                                                                                                                                                                     Score 16.8; D
Pred. No. 1.2e
0; Mismatches
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                                                                                                                                                                                                               DB 4;
                                                                                                                                                                                                               Length 129327;
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Result
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Maximum Match 100%
Listing first 45 s
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                              Score
    16.8
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seq length: 2000000000
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1: /cgn2_6/ptcdata/2/pubpna/PCT_NEW_PUB.seq:*

2: /cgn2_6/ptcdata/2/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptcdata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptcdata/2/pubpna/US07_NEW_PUB.seq:*

9: /cgn2_6/ptcdata/2/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptcdata/2/pubpna/PCTUS_PUBCOMB.seq:*
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                                                                                                                                                                                                                                                                                      Query
                                                                                                                                                                                                                                                                                                                                                                                                                                       / Ggn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq: *
/ Ggn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09B_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10D_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10D_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq: *
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Copyright (c) 1993 - 2005 Compugen Ltd.
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11 US-09-909-317-2
11 US-09-909-317-5
18 US-10-322-281-560
18 US-10-719-993-16987
18 US-10-719-993-16988
10 US-09-764-891-9678
10 US-08-781-986A-1722
17 US-10-329-624-1722
17 US-10-329-624-1722
18 US-10-027-632-218409
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2399.695 Million cell updates/sec
Sequence 2, Appli
Sequence 560, App
Sequence 560, App
Sequence 16987, A
Sequence 16988, A
Sequence 9678, Ap
Sequence 1722, Ap
Sequence 1722, Ap
Sequence 218409,
Sequence 218409,
Sequence 218409,
Sequence 6765, Ap
                                                                                                                                                                                                                                                           Description
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ALIGNMENTS

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RESULT 2
US-09-909-317-5/c
; Sequence 5, Application US/09909317
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                                                                                                                                                                                                                                                                             ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-909-317-2
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                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Betty P. Teao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: 18810-82152
                                                                                                                                                                                         Query Match 100.0%; Score 20;
Best Local Similarity 100.0%; Pred. No.
Matches 20; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/09909317 Publication No. US20040152075A1
                                                                                                                                  1 AAATTGTGGTAATGACTGCA 20
                                                                                                          AAATTGTGGTAATGACTGCA
                                                                                                          20
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GENERAL INFORMATION:

Publication No. US20040152075A1

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CURRENT APPLICATION NUMBER: US/10/322,281
CURRENT FILING DATE: 2002-12-17
NUMBER OF SEQ ID NOS: 866
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 560
LENGTH: 68732
TYPE: DNA
ORGANISM: Homb sapiens
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US-10-719-993-16987/c
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APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
ITILE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEG ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
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                                                                                                                                                         Sequence 16987, Application US/10719993
Publication No. US20040265849A1
GENERAL INFORMATION:
          APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: ALZHEIMER'S DISEASE, METHODS OF DETECTION AND USES
FILE REFERENCE: CLO01496
CURRENT APPLICATION NUMBER: US/10/719,993
CURRENT FILING DATE: 2003-11-24
NUMBER OF SEQ ID NOS: 55342
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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Best Local S
Matches 20
        SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: David W. Morris
APPLICANT: Make S. Malandro
TITLE OF INVENTION: Novel Compositions and Methods in Cancer
FILE REFERENCE: 529452001000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(68732)
OTHER INFORMATION: n = A,T,C or G
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TYPE: DNA
ORGANISM: Homo sapiens
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Local Similarity 100.0%;
hes 20; Conservative 0
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FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                            90.0%; Score 18; DB 18; 100.0%; Pred. No. 2.2e+02
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Pred. No. 13;
0; Mismatches
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hes 0;
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NUMBER OF SEQ ID NOS: 55342

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1698
LENGTH: 201
TYPE: DNA
CRGANISM: Homo sapiens
US-10-719-993-16988
                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/764,891
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper
NUMBER OF SEQ ID NOS: 10231
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 9678
LENGTH: 256
TYPE: DNA
ORGANIZM: Homo sapiens
US-09-764-891-9678
RESULT 7
US-08-781-986A-1722/c
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US-09-764-891-9678
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US-10-719-93-16988/c
; Sequence 16988, Application US/10719993
; Publication No. US20040265849A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9678, Application US/09764891 Publication No. US20030077808A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 84.0%;
Best Local Similarity 90.0%;
Matches 18; Conservative
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: GENETIC POLYMORPHISMS
TITLE OF INVENTION: ALZHEIMER'S DISEASE,
FILE REFERENCE: CLOO1496
CURRENT APPLICATION NUMBER: US/10/719,993
CURRENT FILING DATE: 2003-11-24
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LENGTH: 201
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                                                                                        1 AAATTGTGGTAATGACTGCA 20
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                                                                                                                                 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 AAATTGTGGTAATGACTGCA 20
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                                                               AAATTGTTAATGACTACA 105
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GENETIC POLYMORPHISMS ASSOCIATED WITH
ALZHEIMER'S DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                               84.0%;
90.0%;
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Pred. No. 3.5e+02;
0; Mismatches 2;
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Pred. No. 3.4e+02;
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Pred. No. 3
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                                                                                                                                                              Length 256;
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                               Sequence 1722, Application US/10329624
Publication No. US20040043037A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1722, Application US/08781986A Publication No. US20030054436A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 325 base pairs
                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Benson, Bob
REGISTATION NUMBER: 30,446
REFERENCEJOCKET NUMBER: PB:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: HP Vectra 48
OPERATING SYSTEM: MSDO:
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ALILE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Charles Kunsch
TITLE OF INVENTION: Staph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 9410 Key
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome ScientsTEET: 9410 Key West Avenue CITY: Rockville
STATE: Maryland
                                                                                                                                                                                              Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                                                                                                                                                                                                                                                                                                              APPLICANT: Charles Kunsch
Gil H. Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               57
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18; Conserv
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                                                                       ZIP: 20850
                                                                                            COUNTRY: USA
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                                                                                                                                                                                                                                                                      Patrick S. Dillon
Craig A. Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      84.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MSDOS version 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US/08/781,986A
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                                                                                                                                                              Genome Sciences, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PB248PP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8; DB 8;
Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 325;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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RESULT 9
US-10-027-632-218409/c
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                                                                                                                                                US-10-027-632-218409
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PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR TILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR PPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR PRIOR DATE: 2000-03-29
PRIOR PRIOR DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-09-28
PRIOR PILING DATE: 1999-09-29
PRIOR PILING DATE: 1999-09-29
PRIOR PILING DATE: 1999-09-29
PRIOR PRIOR PRIOR PRIO
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Publication No. US20020198371A1
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                  SEQ ID NO 218409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/027,632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (240) 314-1224
TELEFAX: (301) 309-8439
INFORMATION FOR SEQ ID NO: 1722:
                                                                                                                                                                          ORGANISM: Human
                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                 LENGTH: 663
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APPLICATION NUMBER: US/10/329,624
FILING DATE: 27-Dec-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/956,171
FILING DATE: October 20, 1997
APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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REFERENCE/DOCKET NUMBER: PB248P1D1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 1722:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57 AACGTGTGGTAATGACTGCA 38
   18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 84.0%;
Similarity 90.0%;
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STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Mark J. Hyman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 325 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
Conservative
                                                                                                                                                                                                                                                                                                                           for Windows Version 4.0
                                 90.0%;
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Score 16.8; D
Pred. No. 4.1e
0; Mismatches
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Pred. No. 3.6e+02;
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                                     4.1e+02;
                                                                      DB 13;
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                                                               Length 663;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 325;
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RESULT 10 US-10-027-632-218409/c 유 성

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AAATTGTGGTAATGACTGCA 20

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RESULT 11
US-10-425-114-6765
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                                                                                                                                                         NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 6765
LENGTH: 1386
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  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 6765, Application US/10425114
Publication No. US20040034888A1
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Best Local Similarity
                                                                                                                                                                                                                         APPLICANT: Zhou, Yihua
APPLICANT: Koyalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313) B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Liu, Jingdong APPLICANT: Zhou, Yihua
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/167,363 PRIOR FILING DATE: 1999-11-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
                                                                                                            TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Human
                                                               OTHER INFORMATION: Clone ID: 700605169_FLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             193 AAAATGTGGTAATGACTGTA 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
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  84.0%;
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  Score 16.8;
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Pred. No. 4.1e+02;
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  DB 17;
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Length 1386;
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RESULT 13
US-08-781-986A-38
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APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223)B
FILE REFERENCE: 38-21 (53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 53048
LENGTH: 2591
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APPLICANT: La Rosa Thomas
APPLICANT: Kovalic David
APPLICANT: Zhou Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 38, Application US/08781986A
publication No. US20030054436A1
GENERAL INFORMATION:
APPLICANT: Charles Kunsch
TITLE OF INVENTION: Staphylococcus
NUMBER OF SEQUENCES: 5255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 18; Conserv
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Matches 18; Conserv
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ATTORNEY/AGENT INFORMATION:
NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PB24
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Glycine max
FEATURE:
                                                                                                                                                                                            OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/781,986A
FILING DATE:
                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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9410 Key West Avenue
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                                                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Staphylococcus aureus Polynucleotides and Sequences 5255
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                                         PB248PP
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Pred. No. 5e
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); Mismatches
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5e+02;
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TELEFAX: (301) 309-8512 INFORMATION FOR SEQ ID NO:

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US-08-781-986A-38
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US-10-329-624-38
                                                                                                   US-10-329-624-38
Query Match
Best Local Similarity 90.0
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Matches
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Publication No. US20040043037A1
GENERAL INFORMATION:
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                                                                                                                  TELEFAX: (301) 309-8439
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 23439 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 38:
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APPLICATION NUMBER: US/10/329,624
FILING DATE: 27-Dec-2002
PRIOR APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: 08/956,171
FILING DATE: October 20, 1997
APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT IMFORMATION:

REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB248P1D1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (240) 314-1224
TELEPHONE: (240) 314-1224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 23439 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Charles Kunsch
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18; Conservative
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Craig A. Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gil H. Choi
               84.0%; Score 16.8; DB 17; Length 23439; 90.0%; Pred. No. 7.1e+02; rative 0; Mismatches 2; Indels 0;
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90.0%; Pred. No. 7.1e+02;
tive 0; Mismatches 2;
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APPLICANT: Nicholas M. Dean
APPLICANT: C. Frank Bennett
APPLICANT: C. Frank Bennett
APPLICANT: Kenneth W. Dobie
ITITE OF INVENTION: ANTISENSE MODULATION OF PPP3CB EXPRESSION
FILE REFERENCE: PTS-0028
CURRENT APPLICATION NUMBER: US/10/210,723
CURRENT FILING DATE: 2002-07-31
NUMBER OF SEQ ID NOS: 141
SEQ ID NO 13
LENGTH: 70000
TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                          ; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 63612-63711
; OTHER INFORMATION: n = A,T,C
US-10-210-723-13
Search completed: February 28, 2005, 06:52:29 Job time: 53.3851 secs
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                                                                                                                                        Matches
                                                                                                                                                         Query Match
Best Local Similarity
                                                                      39997 AAACTGTGGTAATGATTGCA 39978
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                                                                                                                                        Conservative
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Result
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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35.4
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Match Length
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfIles1.seq:*
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Maximum Match 100%
Listing first 45 summaries
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      US-09-949-016-203106
US-09-943-016-17468
US-09-943-016-13589
US-09-949-016-131388
US-09-949-016-22412
US-09-949-016-22411
US-09-949-016-154217
US-09-949-016-154217
US-09-949-016-11957
US-09-949-016-142943
US-09-949-016-142943
US-09-949-016-142943
US-09-949-016-13256
US-09-949-016-13110
US-09-949-016-15779
US-09-949-016-15779
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US-09-949-016-15779
US-09-949-016-15779
US-09-949-016-15779
US-09-949-016-15779
US-09-949-016-13110
US-09-949-016-15779
US-09-949-016-13110
US-09-949-016-13439
                                                                                                                                                                                                                                                                                                                               SUMMARIES
Sequence 203106, Sequence 13748, A Sequence 13709, A Sequence 11808, A Sequence 13808, A Sequence 13417, A Sequence 154217, Sequence 154217, Sequence 16064, A Sequence 16064, A Sequence 142943, Sequence 142944, Sequence 13256, A Sequence 11894, A Sequence 13779, A Sequence 51612, Appli Sequence 51612, Appli Sequence 51612, Appli Sequence 51613, Appli Sequence 51613, Appli Sequence 1331, A Sequence 13439, A Sequence 14349, A Sequence 14349, A Sequence 14349, A Sequence 14350, A
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SOFTWARE: FastSEQ for Windows SEQ ID NO 17468

Version

PRIOR FILING DATE: 2000-09-08 NUMBER OF SEQ ID NOS: 207012

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                                                                                                                                                                                                                                                                                                                             RESULT 4
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961

FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION UNMBER: US 60/122,487
PRIOR PILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                         Sequence 17009, Application US/09949016

Patent No. 6812339

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
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SEQ ID NO 13589
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: buclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
             PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 182
TYPE: DNA
ORGANISM: Homo sapiens
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LOCATION: (1)...(88906)
OTHER INFORMATION: n = A,T,C
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ORGANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                  TCTCTTTCTTTACACACACACACACACACACACACAAAATATCT 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       72.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     70.6%;
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Pred. No. 0.0022;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 36; DB 4;
Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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SEQUENCE 13388, Application US/09949016

Patent No. 6812339

PATENT INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 
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US-09-949-016-11808/c
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 17009
LENGTH: 205163
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Best Local Similarity
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(636591)
OTHER INFORMATION: n = A,T,C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Human
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97.3%;
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Pred. No. 0.014;
0; Mismatches 10; Indels 0
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OF DETECTION AND USES
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US-09-949-016-22414/c
; Sequence 22414, Application US/09949016
; Patent No. 6812339
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SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 22412
LENGTH: 601
TYPE: DNA
ORGANISM: Human
US-09-949-016-22412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-949-016-22412/c; Sequence 22412; Appl: Patent No. 6812339; GENERAL INFORMATION:
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; LOCATION: (1)...(636591)
; LOCHER INFORMATION: n = A,T,C
US-09-949-016-13388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 22414
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                                                           PRIOR APPLICATION NUMBER: 60/241.755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
                                           NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: CL001307
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Local Similarity 80.4%;
les 41; Conservative
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Pred. No. 0.0041;
0; Mismatches 7;
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Pred. No. 0.014;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 601;
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                                      US-09-949-016-154219
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILLING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6812339
GENERAL INFORMATION:
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                SEQ ID NO 154219
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 154219,
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Query Match
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                                                                                                                                                                APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
                                                                                                                                NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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                                                          ORGANISM: Human
                                                                            TYPE: DNA
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                                                                                            LENGTH: 601
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Similarity 84.8%;
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                                                                                                                                    for Windows Version
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  68.2%;
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  Score 34.8;
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Pred. No. 0.0041;
0; Mismatches
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Pred. No. 0.0041;
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Length 601;
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PRIOR APPLICATION NUMBER: 60/241.755
PRIOR APPLICATION NUMBER: 60/241.755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11957
LENGTH.
                                                                                                                                                                                                                                                                    GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR PILING DATE: 2000-10-03

PRIOR PILING DATE: 2000-0-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08
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US-09-949-016-11957
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US-09-949-016-11957
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                                                                Query Match 68.2
Best Local Similarity 84.8
Matches 39; Conservative
                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16064
LENGTH: 70131
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                                                                                                                                                   ORGANISM: Human
-09-949-016-16064
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 11957, A
Patent No. 681233
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CURRENT FILING DATE: 2000-04-14
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39; Conservative
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ATCTCTTTCTTTACACACACACACACACACACACACAAAATATCTG 49
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                                                             Score 34.8; DB 4; Length 70131;
Pred. No. 0.01;
0; Mismatches 7; Indels 0;
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OF DETECTION
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION F
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: U9/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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US-09-949-016-142944
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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US-09-949-016-142943
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                                                                                                    Query Match
Best Local Similarity
                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 142943
LENGTH: 601
TYPE: DNA
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Best Local
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GENERAL INFORMATION:
APPLICANT: VENTER, J.
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                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                   LENGTH: 601
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214
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                                     Conservative
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                                                                                                    67.8%;
90.2%;
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Pred. No. 0.0048;
0; Mismatches 4;
                                                                                               Score 34.6; DB 4; Length 601; Pred. No. 0.0048;
                                                                                Mismatches
                                                                                4.
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RESULT 15 US-09-949-016-142945

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Fatent No. 681239

[SAMERAL IMPORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
ITILE OF INVENTION: POLYMORPHISMS.
ITILE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF ILLE REFERENCE: CLOO1307

CURRENT PELLICATION UNMERE: US/09/949,016
[CURRENT FILING DATE: 2000-04-14]
PRIOR APPLICATION UNMERE: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,756
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR REPLICATION NUMBER: 60/237,766
PRIOR APPLICATION NUMBER: 60/237,766
PRIOR APPLICATION NUMBER: 60/237,498
PRIOR FILING DATE: 2000-10-90-08
IMMBER OF SEQ ID NOS: 207012
SEQ ID NO 149945

LENGTH: 601
TYPE: DIN
ORGANISM: Human
US-09-949-016-142945

Query Match
Best Local Similarity 90.2%; Pred. No. 0.0048;
Matches 37; Conservative 9; Mismatches 4; Indels 0; Gaps 0;
DETECTION OF CONTINUED OF CONTINU
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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Copyright (c) 1993 - 2005 Compugen Ltd.
/Ggn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US09B_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US09B_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*
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/Ggn2_6/ptodata/2/pubpna/US10D_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US10D_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US10D_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
/Ggn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
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/Ggn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

c c 10 5 8	0.0	000	Result
36.8 36.8 36.8	38.6	51 40.4 40.4	Score
72.5 72.2 72.2 72.2	79.2 75.7 73.3	100.0 79.2 79.2	% Query Match
72.5 621 72.2 104245 72.2 104245 72.2 104245 72.2 170245	10619	2085 10619 10619	% Query Match Length DB
18 17 17 18	17 18	114	
US-10-357-930-55243 US-10-160-807-4 US-10-655-847-4 US-10-717-597-322	US-10-240-589C-2 US-10-674-124A-3039 US-10-674-124A-10920	US-09-909-317-5 US-10-239-676-2 US-10-311-455-44	SUMMARIES ID
Sequence 55243, A Sequence 4, Appli Sequence 4, Appli Sequence 322, App	Sequence 2, Appli Sequence 3039, Ap Sequence 10920, A	Sequence 5, Appli Sequence 2, Appli Sequence 44, Appl	Description

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4 4.	44	3	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12
34.8			•	•	٠	٠	•	•	٠	٠	•	٠	•				٠	٠	•		٠	•	•		•	•	•	٠	٠	٠	36	36	36.2
68.2	68.2	•	•	•	•	•	•	•	•	•	•	•	٠	٠	٠	٠	٠	•	•		69.0	•	٠	•	•	•	•	•	69.8	•	•	•	71.0
112241	96649	96649	23503	598	598	457	448	357	286	252	250	188	134	14277	727	727	727	727	727	727	570	570	393	289	143	124	576	576	58723	143	657	657	419
18	17	10	18	17	13	18	18	18	18	18	18	18	18	18	17	17	17	13	13	13	17	13	18	18	18	18	17	13	13	18	17	13	18
-10-322-281- ALIGNMENTS	10-633-913	-09-956-712-	-10-322-281-	-10-027-6	-10-027-632-	-10-674-1	US-10-674-124A-3935	-10-674-124A-584	-10-674-124A-1052	-10-674-124A-190	US-10-674-124A-10989		-10-674-124A-943	-10-715-066-	-10-027	-10-027-632-14946	-10-027	-10-027-632-14946	-10-027-632-14946	-10-027-632-14945	027	-10-027-632-12816	-10-674-124A-2524	-10-674-124A-2322	-10-674-124A-2525	-10-674-124A-1202	-10-027	US-10-027-632-282410	-10-087-192-4	US-10-674-124A-12309	US-10-027-632-259460	-10-027-632-25946	US-10-674-124A-18213
656	Sequence 10, Appl	10,	667	e 180	18056	e 17215,	3935,	5849,	1052	19062,	e 10989,	6272,	e 9430,	3, Ap	e 149	Œ	ø	ø	O	æ	Sequence 128160,	æ	e 25249,	O	equence 25250,	e 12025,	equence	28241	Œ	æ	e 25946	Ø N	Sequence 18213, A

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; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-10-311-455-44
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Sequence 44, Application US/10311455
Publication No. US20030143606A1
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                                                                                Matches
                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-09-01; NUMBER OF SEQ ID NOS: 228; SEQ ID NO 2; LENGTH: 10619; TYPE: DNA; ORGANISM: Artificial Sequence; PEATURE:
                                                                                                                                                                                                                                 SEQ ID NO 44
LENGTH: 10619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 97.6
41; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION
                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 2424
                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/311,455
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: PCT/EP01/07537
PRIOR FILING DATE: 2001-07-02
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: PIÉPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Determ
TITLE OF INVENTION: cytosine methylation
FILE REFERENCE: 5013.1014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, C
                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DE 10019058.8

DE 10019173.8

DE 1002529.7

DE 10043826.1

PRIOR FILLING DATE: 2001-04-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/239,676
CURRENT FILING DATE: 2002-09-24
PRIOR APPLICATION NUMBER: PCT/EP01/03968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with Gene Regulation
FILE REFERENCE: 5013.1003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                              Local Similarity
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decarciónintentracacadadadadadadadadadadada 663
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                                                                                          79.2%;
97.6%;
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                                                                                     Score 40.4; DB 15
Pred. No. 0.00034;
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Pred. No. 0.00034;
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                                                                                                       DB 15;
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                                                                     Indels
                                                                                                    Length 10619;
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CURRENT APPLICATION NUMBER: US/10/240,589C
CURRENT FILING DATE: 2003-09-02
PRIOR APPLICATION NUMBER: PCT/EP01/03972
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR PILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-06-30
PRIOR PILING DATE: 2000-06-30
PRIOR PILING DATE: 2000-06-30
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                                                                                              SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2002-10-02
PRIOR APPLICATION NUMBER: PCT/EP01/03973
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR PILING DATE: 2000-04-06
PRIOR PILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR PILING DATE: 2000-06-30
PRIOR PILING DATE: 2000-09-01
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/10240589C
Publication No. US20040076956A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with
FILE REFERENCE: 5013.1008
                                                                                                                PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 148
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Best Local Similarity
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SEQ ID NO 2
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/10240453 Publication No. US20030148326A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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FILE REFERENCE: 5013.1009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Transcription
TITLE OF INVENTION: by Means of Assessing the Methylation St
TITLE OF INVENTION: With DNA TRANSCription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/240,453
CURRENT FILING DATE: 2002-10-02
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97.6%;
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Pred. No. 0.00034;
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                                                                                                                                                                                                                         US-10-674-124A-10920
                                                                                                                                                                                                                                                      RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/674,124A
CURRENT FILING DATE: 2003-09-26
PRIOR APPLICATION NUMBER: 10/257,511
PRIOR FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: JP2000-112699
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-327516
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                                                                                                                                           Sequence 10920, Application US/10674124A Publication No. US20040197797A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
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SEQ ID NO 3039
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       APPLICANT: INOKO, Hidetoshi
APPLICANT: TAMIYA, Gen
TITLE OF INVENTION: GENE MAPPING METHOD USING MICROSATELLITE
TITLE OF INVENTION: GENETIC POLYMORPHISM MARKERS
FILE REFERENCE: ORLN-003CIP
FILE REFERENCE: ORLN-003CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: INOKO, Hidetoshi
APPLICANT: TAMIYA, Gen
TITLE OF INVENTION: GENE MAPPING METHOD USING MICROSATELLITE
TITLE OF INVENTION: GENETIC POLYMORPHISM MARKERS
FILE REFERENCE: ORIN-003CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: JP2002-383869 PRIOR FILING DATE: 2002-12-09
CURRENT APPLICATION NUMBER: US/10/674,124A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PEATURE:
OTHER INFORMATION: Distance between 3'-terminus of neighbour sequence of OTHER INFORMATION: sequence listing upward to telomere on chrosomal short arm and OTHER INFORMATION: 5'-terminus of this base sequence : 18142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Distance between a terminus base of telomere on OTHER INFORMATION: chromosomal short arm and 5'-terminus of this base OTHER INFORMATION: sequence: 84667136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Located on chromosome 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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41; Conserv
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91.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                            Score 38.6; DB 18 Pred. No. 0.00079; 0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 40.4;
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 400;
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                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                           APPLICANT: Schlegel. Robert
APPLICANT: Schlegel, Wilson
APPLICANT: Mondhan, John
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS,
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PRI
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PRI
TITLE OF INVENTION: UNMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: 09/785,276
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/189,862
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILING DATE: 2000-03-16
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILING DATE: 2000-05-25
PRIOR FILING DATE: 2000-05-25
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PRIOR APPLICATION NUMBER: 10/257,511
PRIOR FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: JP2000-112699
PRIOR FILING DATE: 2000-04-13
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR FILING DATE: 2002-09-28
PRIOR FILING DATE: 2002-09-28
                                                                      SOFTWARE: FastSEQ for Windows Version SEQ ID NO 55243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 55243, Application US/10357930 Publication No. US20040259086A1
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SEQ ID NO 10920
                                                                                                                                             PRIOR APPLICATION NUMBER: 60/211,314
PRIOR FILING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR FILING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: JP2002-383869
PRIOR FILING DATE: 2002-12-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Distance between 3'-terminus of neighbour sequence of OTHER INFORMATION: sequence listing upward to telomere on chrosomal short arm and OTHER INFORMATION: 5'-terminus of this base sequence: 107348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Distance between a terminus base of telomere on OTHER INFORMATION: chromosomal short arm and 5'-terminus of this base OTHER INFORMATION: sequence : 131608898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 154
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Located on chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: D6S988
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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                                               ENGTH: 621
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 TCTCTTTCTTTACACACACACACACACACACACACAAAATATCTGAT 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ID NOS: 27110
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87.2%;
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Length 154;

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Gaps

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AND METHODS

FOR THERAPY OF

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LENGTH: 104245

TYPE: DNA

CRGANISM: Homo sapiens

FEATURE:
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                                                                    Matches
                                                                               Best Local
                                                                                              Query Match
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Publication No. US20040063129A1
                                                                                                                              -10-655-847-4
                                                                                                                                                                                                          APPLICANT: William Gaarde
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
CURRENT FAPLICATION NUMBER: US/10/655,847
CURRENT FLING DATE: 2003-09-05
PRIOR APPLICATION NUMBER: US/10/160,807
PRIOR FILING DATE: 2003-09-05
NUMBER: OF SEQ ID NOS: 296
NUMBER OF SEQ ID NOS: 296
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LENGTH: 104245
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Publication No. US20030224514A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/160,807
CURRENT FILING DATE: 2002-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: n = A,T,C or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90002
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                                                                   41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41;
                                                                              Similarity
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TCTCTTTCTTTACACACACACACACACACACACACAAAATATCTG 49
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                                                                              72.2%;
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                                                           Score 36.8; DB 17;
Pred. No. 0.0093;
0; Mismatches 7;
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                                                                                         Length 104245;
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CURRENT FILING DATE: 2003-09-26
PRIOR APPLICATION NUMBER: 10/257,511
PRIOR FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR APPLICATION NUMBER: JP2000-112699
PRIOR FILING DATE: 2000-04-13
PRIOR PLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-383869
PRIOR PILING DATE: 2002-09-28
PRIOR PILING DATE: 2002-12-09
PRIOR FILING DATE: 2002-12-09
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US-10-717-597-322
                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 27110
SEQ ID NO 18213
LENGTH: 419
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Matches
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Publication No. US20040197797A1
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LENGTH: 170245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: ORIN-003CIP
CURRENT APPLICATION NUMBER: US/10/674,124A
CURRENT FILING DATE: 2003-09-26
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Publication No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: TAMIYA, Gen
TITUE OF INVENTION: GENE MAPPING METHOD USING MICROSATELLITE
TITUE OF INVENTION: GENETIC POLYMORPHISM MARKERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: INOKO, APPLICANT: TAMIY
                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                            TYPE: DNA
OTHER INFORMATION: Distance between a terminus base of telomere on OTHER INFORMATION: chromosomal short arm and 5'-terminus of this base
                                                                  OTHER INFORMATION: Located on chromosome 12
                                                                                                          FEATURE:
OTHER INFORMATION: AC004802.1_36841
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SOFTWARE: PatentIn version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/459,782
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: US 60/427,982
PRIOR FILING DATE: 2002-11-21
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TITLE OF INVENTION: METHODS FOR DIAGNOSING RCC AND OTHER SOLID TUMORS
FILE REFERENCE: AM101080L
CURRENT APPLICATION NUMBER: US/10/717,597
CURRENT FILING DATE: 2003-11-21
                                                       FEATURE
                                                                                               FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Slonim, Donna K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dorner, Andrew J.
Trepicchio, William L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Twine, Natalie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Burczynski, Michael E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10717597 b. US20040110221A1
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85.4%;
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                                                                                                                                                                                                                                                                                      RESULT 14
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US-10-027-632-259460
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GENERAL INFORMATION:

APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide

TITLE OF INVENTION: Polymorphisms in the Human Genome.

FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US/10/027,632

CURRENT FILING DATE: 2002-04-30

PRIOR APPLICATION NUMBER: US 60/218,006

PRIOR FILING DATE: 2000-07-12

PRIOR APPLICATION NUMBER: US 60/198,676

PRIOR FILING DATE: 2000-04-20
                                                                                                                                                                                                                        Sequence 259460, Application US/10027632 Publication No. US20030204075A9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 259460
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Sequence 259460, Application US/10027632
Publication No. US20020198371A1
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
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APPLICATION NUMBER: US 60/146,002
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Pred. No. 0.00
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PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR FILING DATE: 1999-11-23

PRIOR APPLICATION NUMBER: US 60/156,358

PRIOR FILING DATE: 1999-09-28

PRIOR FILING DATE: 1999-09-28

PRIOR APPLICATION NUMBER: US 60/146,002

PRIOR FILING DATE: 1999-08-09

PRIOR FILING DATE: 1999-08-09
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SEQ ID NO 12309
LENGTH: 143
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR FILING DATE: 2002-09-28
PRIOR APPLICATION NUMBER: JP2002-383869
PRIOR FILING DATE: 2002-12-09
NUMBER FILING DATE: 2002-12-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/674,124A
CURRENT FILING DATE: 2003-09-26
PRIOR APPLICATION NUMBER: 10/257,511
PRIOR FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR PRIOR PILING DATE: 2000-10-30
PRIOR PILING DATE: 2000-04-13
PRIOR FILING DATE: 2000-04-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: GENE MAPPING METHOD USING MICROSATELLITE TITLE OF INVENTION: GENETIC POLYMORPHISM MARKERS FILE REFERENCE: ORIN-003CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                   OTHER INFORMATION: Distance between 3'-terminus of neighbour sequence of OTHER INFORMATION: sequence listing upward to telomere on chrosomal shor OTHER INFORMATION: 5'-terminus of this base sequence: 39232
                                                                                                                                                                              OTHER INFORMATION: Distance between a terminus base of telomere of the information: chromosomal short arm and 5'-terminus of this other information: sequence: 81430512
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                                                                                                                                                                   FEATURE:
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                                                                                                                                                                                                                                                                    OTHER INFORMATION: Located on chromosome
                                                                                                                                                                                                                                                                                            FEATURE:
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US-09-517-4678-3
US-08-044-618-5
US-08-05-248D-24
US-08-05-05-05-048D-46
US-09-596-248D-46
US-09-596-016-14854
US-09-949-016-13423
US-09-949-016-150-1609
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Sequence 1,8058,
Sequence 16047, A
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SULT 2 -08-044-618-5 -08-044-618-5 Sequence 5, A Patent No. 54 GENERAL INFC APPLICANT: TITLE OF I TITLE OF I TITLE OF I TITLE OF I	y 1188 y 188 y 194 y 194 y 194 b 10 y 194 y 200 b 22	O9-517-467B eequence 3, p atent No. 64 ENERAL INFOR APPLICANT: TITLE OF INF TOWNER OF S EN IN OF S LENGTH: 366 TYPE: DNA ORGANISM: H FEATURE: NAME/KEY: C LOCATION: (109-517-467B		4 4 4 4 4 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
NV NV NV	5 8 5 8 7 5	467B-3 3, Appl 3, Appl 10.64516 INFORMATI VI: IAN VI: IAN VI: IAN FILING FILING DA FIL		44444444444444444444444444444444444444
licatio 605 AMIION: SMULSON ENTION: ENTION: ENTION:	Conser Conser Conser JTTTCTA JTTTCTA JTTTCTA JTTTCTTCA AGTCTTCA AGTCTTCA AGTCTTCA AGTCTTCA	Hical		
N. ON POP	Ty 100 control to the	RESULT 1 US-09-517-467B-3 Sequence 3, Application US/0951746 Patent No. 6451602 GENERAL IMPORMATION: APPLICANT: IAN PODOFF APPLICANT: IAN PODOFF APPLICANT: IAN PODOFF APPLICANT: IAN PODOFF APPLICANT APPLICATION UMMER: US/09 FILE REFERENCE: RTS-0150 CURRENT APPLICATION UMMER: US/09 CURRENT FILING DATE: 2001-03-02 PRIOR APPLICATION UMMER: 09/517, PRIOR FILING DATE: 2000-03-02 PRIOR FILING DATE: 2000-03-02 NUMBER OF SEQ ID NOS: 345 SEQ ID NO 3 LENGTH: 3660 TYPE: DNA ORGANISM: Homo sapiens FEATURE: NAME/KEY: CDS LOCATION: (160)(3204) US-09-517-467B-3		601 2252 2273 8907 113042 113142 113142 152145 177251 1638 16738 16738 16738 16738 16738
'0804' 'HOD ('HOD ('YMOR'	GGCTC	0951 NSE 1-03-09/E		444444444444
US/08044618 MARK METHOD OR DETECTING A PREDISPOSITION CANCER BY THE USED OF RESTRICTION FR. POLYMORPHISM OF THE GENE FOR THE HUM POLYMERASE	imilarity 100.0%; Pred. No. 6e-59; Conservative 0; Mismatches 0; Indels 0; Gaps GTGTTTCTAGGTCGTGGGGTCGGGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAAGGATGGC	7467B MODULATION OF PARP EXE 09/517,467 17,467	ALIGNMENTS	US-09-949-016-15193 US-09-949-016-1519 US-09-949-016-11761 US-09-949-016-12343 US-09-949-016-12343 US-09-949-016-12371 US-09-949-016-12371 US-09-949-016-12371 US-09-949-016-12371 US-09-949-016-12468 US-09-949-016-12468 US-09-949-016-12168 US-09-949-016-12168 US-09-949-016-12168 US-09-949-016-12168 US-09-949-016-12168 US-09-949-016-12168 US-09-949-016-12468 US-09-949-016-13527 US-09-949-016-15527 US-09-949-016-15527 US-09-949-016-15527 US-09-949-016-15527
TO AGMENT	Indels 0; G GCATCATGGTGAGA CCATCATGGTGAGA	NOI		Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence
LENGTH (ADP-RIBOSE)	Gaps 0; ATGGC 1947 ATGGC 164 ATGCAA 2007 3 2062			161293, 4519, Ap 119, Appl 119, Appl 119, Appl 112343, A 112343, A 112343, A 113845, A 113845, A 113847, A 1158479, 810, Appl 810, Appl 810, Appl 14678, A 14948, A 14948, A

NUMBER OF SEQUENCES:

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RESULT 3
US-09-596-248D-24
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REFERENCE/DOCKET NUMBER: 0654.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)466-0800
TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO: 5:
SECHIENCE CENTRALIES.
          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 24
LENGTH: 3045
                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                               APPLICANT: DeMaggio, Anthony J
APPLICANT: Goldman, Phyllis S
APPLICANT: McBlilgott, David L
TITLE OF INVENTION: Human Poly(ADP-Ribose) Polymerase 2 Materials and
TITLE OF INVENTION: Methods
FILE REFERENCE: 27866/36544
CURRENT APPLICATION NUMBER: US/09/596,248D
CURRENT FILLING DATE: 2000-06-16
                                                                                                                                                                                                                                APPLICANT: Christenson, Erik
APPLICANT: DeMaggio, Anthony
APPLICANT: Goldman, Phyllis
APPLICANT: McElligott, David
                                                                         PRIOR APPLICATION NUMBER: 60/139,543
PRIOR FILING DATE: 1999-06-16
NUMBER OF SEQ ID NOS: 68
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/044,618
TYPE: DNA
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APPLICATION NUMBER: US/07/257,696
FILING DATE: 14-OCT-1988
ATTORNEY/AGENT INFORMATION:
NAME: FOX, SAMUEL L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: 1
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LENGTH: 3747 base pairs
TYPE: NUCLEIC ACID
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STATE: L.
To: 20036
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CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                          24, Application US/09596248D
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E: DNA
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100.0%; Pred. No. 6e-59;
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; FEATURE:
; NAME/KEY: Coding S
; LOCATION: 96...313
; OTHER INFORMATION:
US-08-860-886-1
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US-09-596-248D-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 4
                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAURA A
REGISTRATION NUMBER: 30,742
REFERRINCE/DOCKET NUMBER: 8484
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
Query Match 3.5%;
Best Local Similarity 98.9%;
Matches 173; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 5.8%; So Best Local Similarity 100.0%; E Matches 120; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1,
                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: Windows SOFTWARE: FastSEQ for Windows Version 2. CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/860,886 FILING DATE: 03-OCT-1997 CLASSIFICATION: DATA: APPLICATION UMBER: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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LOCATION: (1)..(3045)
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                  LENGTH: 3792 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Pennie & Edmonds, LLP STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61
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Zur Hausen, Harald
Jan-Heiner, Kupper
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                                                                                                     Coding Sequence 96...3134
                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VECTORS
Score 73; DB 3; Le
Pred. No. 3.1e-19;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 120; DB 4;
Pred. No. 1.6e-37;
                                                                                                                                                                                                                                                                                                                               8484-0028-999
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                                 Length 3792;
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US-09-596-248D-46
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                                                                                                                                                                                                                                                   Sequence 7, Application US/08044618 Patent No. 5449605
GENERAL INFORMATION:
APPLICANT: SMULSON, MARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 46
LENGTH: 3200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Christenson, Erik
APPLICANT: Christenson, Erik
APPLICANT: Demaggio, Anthony J
APPLICANT: Goldman, Phyllis S
APPLICANT: McElligott, David L
TITLE OF INVENTION: Human Poly(ADP-Ribose) Polymerase 2 Materials and
TITLE OF INVENTION: Methods
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Best Local (
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CURRENT APPLICATION NUMBER: US/09/596,248D
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: 60/139,543
PRIOR FILING DATE: 1999-06-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 68
                                                                                                                               TITLE OF INVENTION: MET!
TITLE OF INVENTION: CANUTITLE OF INVENTION: POLIVITILE OF INVENTION: POLIVINGHER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence:hPARP1/hPARP2 OTHER INFORMATION: Fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
STATE: D.C. ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                          STREET: 1220
CITY: Washington
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Local Similarity 100.0%;
                                                                                                                   ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          2003 TGCAAGAAATG 2013
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                                                                                                                                                                                                                                                 SMULSON,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                          METHOD OR DETECTING A PREDISPOSITION TO CANCER BY THE USED OF RESTRICTION FRAGMENT LENGTH POLYMORPHISM OF THE GENE FOR THE HUMAN POLY (ADP-
                                                                                                                                                                            POLYMERASE
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; Pred. No. 1.9e-1
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ches 0;
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US-09-949-016-14854
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                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/231,768
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version SEQ ID NO 14854
LENGTH: 8848
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                                                                                                              Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 5345 base pa
TYPE: NUCLEIC ACID
STRANDENESS: both
TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                              TYPE: DNA
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REGISTRATION NUMBER: 30,353
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)466-0800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/257,696
FILING DATE: 14-OCT-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/044,618
FILING DATE: 19930406
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4278 GGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACAT 4229
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                          770 GGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACAT
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SYSTEM: PC-DOS/MS-DOS
                                                                                      2.4%; Score 50; DB
100.0%; Pred. No. 2.0
tive 0; Mismatches
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3. 9.6e-16;
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2.6e-10;
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                    Sequence 13423, Application US/09949016 Patent No. 6812339
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US-09-949-016-178057
GENERAL INFORMATION
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Best Local :
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Best Local Similarity 100.0%; Pred. No. 1.3e-08;
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 178057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                             ENGTH: 601
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                                                                                                                                                       306 GGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAG 351
                                                                                                                                         302
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
FRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
PRIOR FI
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US-09-949-016-16847/c
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                                        Sequence 15078, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT PILING DATE: 2000-04-14
PRIOR APPLICATION UNMBER: 60/241,755
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
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SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 16847
LENGTH: 98302
TYPE: DNA
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/949,016
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TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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Conservative 0; Mismatches
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Conservative (
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100.0%; Pred. No. 7.3e-09;
ive 0; Mismatches 0;
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Pred. No.
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RESULT 14
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; ORGANISM: Homo sapiens
US-09-513-999C-16098
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PATENT OF INVENTION: Expressed Sequence Tags Patent No. 6783961
FILE REFERENCE: 59 US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
RUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                  Sequence 16110, Application US/09513999C Patent No. 6783961 GENERAL INFORMATION:
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Best Local Similarity
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SOFTWARE: Patent.pm
SEQ ID NO 16098
LENGTH: 73
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags
Patent No. 6783961
FILE REFERENCE: 59.US2.REG
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Best Local Similarity
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 15078
                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                            APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
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PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
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Pred. No. 6.9e-09;
0; Mismatches 0;
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US-09-513-999C-29990
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Search completed: February 28, 2005, 08:24:07 Job time : 377 secs
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LENGTH: 74
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SEQ ID NO 29990
LENGTH: 123
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
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                                                                                                                    Query Match
Best Local Similarity
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GENERAL INFORMATION:
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Best Local (
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CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
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                                                                                                                                                                                                                                                        LOCATION: 18
OTHER INFORMATION: v=a or
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LOCATION: 17
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OTHER INFORMATION: 8=9
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NAME/KEY: misc_feature
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Maximum DB
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seq length: 2000000000
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1: /cgn2_6/ptodata/2/pubpna/US07 PUBCOMB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

6: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

8: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

10: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

11: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*

15: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

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24: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

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Sequence 5, Appli
Sequence 154183,
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Sequence 327, App
Sequence 328, Ap
Sequence 5037, App
Sequence 117, App
Sequence 117, App
Sequence 106, App
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Sequence 316, App
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ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 2085
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-909-317-5
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Publication No. US20040152075A1

GENERAL INFORMATION:

APPLICANT: Betty P. Tsao (Inventor)

APPLICANT: Rita M. Cantor (Inventor)

APPLICANT: Jerome I. Rotter (Inventor)

TITLE OF INVENTION: Genetic Marker Test for Lupus

FILE REFERENCE: 18810-82152
                                                                                                                                   Query Match 100.0%; Score 2085; Best Local Similarity 100.0%; Pred. No. 0; Matches 2085; Conservative 0; Mismatches
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                    61 TTCTCTAAGTAAAACTTTTATTTGTTCCATCATATTTTCCACTTATTCTGTTTTACCTTCA 120
                                                                Mismatches
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	TGCAGGGTCCACCTCGGGCCAATCAACTATATTCCCGAGGCGGGGGCCTGCGCTTCCCGG 1200 	141 141	•
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	ATGAAAACGTCTGTGATGACTATTGCCCACAAAGTCACAGGTACTGCTAATACTC 1020 	961 CTAACATGAAAAC 961 CTAACATGAAAAC	- '
	CACAATTICGAGTAGTGAIGAGCITAATAATATTICGAGTTATCACCAACAACTGTAAA 960 	901 CCACAATTTCGAG	
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RESULT 2
US-10-027-632-154183
US-10-027-632-154183
; Sequence 154183, Application US/10027632
; Publication No. US20020198371A1
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Hodentification and Mapping of Single Nucleotide
; TITLE OF INVENTION: Polymorphisms in the Human Genome
; TITLE OF INVENTION: Polymorphisms in the Human Genome
; TILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; CURRENT FILING DATE: 2002-04-30
; PRIOR APPLICATION NUMBER: US 60/218,006
; PRIOR FILING DATE: 2000-07-12

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RESULT 3
US-10-027-632-154183
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; ORGANISM: Human
US-10-027-632-154183
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APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/193,483
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SOFTWARE: FAStSEQ for Windows Version
SEQ ID NO 154183
LENGTH: 844
                                                                                                                                                                                     Sequence 154183, Application US/10027632 Publication No. US20030204075A9
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Best Local Similarity
Matches 456; Conserv
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Pred. No. 2e-187;
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SEQ ID NO 154183
LENGTH: 844
TYPE: DNA
ORGANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
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Matches 456;
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Best Local Similarity
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                CAGCCCAAGCCCCGCCTCGGCCCCCAGGGCAGCGGCC 1663
                                                          GACTTCCTGGGACAGAACAATCAAAGGGGTGCCCGGGTCCTCCAAAGAGCTACTAGCT
                                                                                GACTTCCTGGGACAGAACAATCAAAGGGGTGGCGCCGGGTCCTCCAAAGAGCTACTAGCT
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Pred. No. 2e-187;
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RESULT 4

US-10-283-975A-327

; Sequence 327, Application US/10283975A
; Publication No. US20040110792A1
; Publication No. US20040110792A1
; Publication No. US20040110792A1
; Publication No. US20040110792A1
; GENERAL INFORMATION: Methods For Assessing and Treating Leukemia FILLR REFERENCE: CDS 293 PCT
; CURRENT APPLICATION NUMBER: US/10/283,975A
; CURRENT FILING DATE: 2002-10-30
; PRIOR APPLICATION NUMBER: 60/340,938
; PRIOR APPLICATION NUMBER: 60/340,938
; PRIOR APPLICATION NUMBER: 60/340,081
; PRIOR APPLICATION NUMBER: 60/340,081
; PRIOR APPLICATION NUMBER: 60/340,081
; PRIOR APPLICATION NUMBER: 60/341,012
; PRIOR FILING DATE: 2001-10-30
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APPLICANT: AZIZ, Natasha
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Zlotnik, Albert
ITILE OF INVENTION: Methods of Diagnosis of Soft Tissue Sarcoma, Compositions &
ITILE OF INVENTION: Methods for Screening for Soft Tissue Sarcoma Modulators
FILE REFERENCE: 0588.2019.NPUSO1
CURRENT APPLICATION NUMBER: US/10/723,860
CURRENT FILING DATE: 2003-11-26
PRIOR FILING DATE: 2002-11-26
ANUMBER OF SEQ ID NOS: 8393
SOFTWARE: Patentin version 3.2
LENGTH: 370
TYPE: DNA
ORGANISM: Homo sapiens
US-10-723-860-2326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2326, Application US/10723860 Publication No. US20040253606A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 175; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version SEQ ID NO 327 LENGTH: 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: HUMAN
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(335)
OTHER INFORMATION: N=any base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PEATURE:

NAME/KEY: misc_feature
LOCATION: (1)...(335)
OTHER INFORMATION:
                                          2008
                                                                                                                                                                                                 1888 GTGTTTCTAGGTCGTGGCGTCGGGCTTCCGGAGCTTTGGCCGGCAGCTAGGGGAGGATGGC 1947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2008 GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 2062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1948 GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAA 2007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1888 GTGTTTCTAGGTCGTGGCGTCGGGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGC 1947
  249
                                                                                     189
                                                                                                                                                                          129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        106 GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGGGGGCGCCTCTTGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 GTGTTTCTAGGTCGTGGCGTCGGGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGC
                          GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 2062
                                                                                                           GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCCGCCTCTTGCAA
GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 303
                                                                                   GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAA
                                                                                                                                                                      GTGTTTCTAGGTCGTGGCGTCGGGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAAGGATGGC 188
                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                         8.4%; Score 175; DB 18; Length 370; 100.0%; Pred. No. 1.3e-74; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8.4%; Score 175; DB 18; 100.0%; Pred. No. 1.3e-74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 335;
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                                                                                                                             2007
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RESULT

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Sequence 5037, Application US/09918995

| Publication No. US20030073623A1
| GENERAL INFORMATION:
| APPLICANT: Hyseq, Inc.
| TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES FILE REFERENCE: 2041-756
| CURRENT APPLICATION NUMBER: US/09/918,995
| CURRENT FILING DATE: 2001-07-30
| PRIOR APPLICATION NUMBER: US/09/235,076
| PRIOR APPLICATION NUMBER: US/09/235,076
| PRIOR FILING DATE: 1999-01-20
| NUMBER OF SEQ ID NOS: 38054
| SOFTWARE: FastSEQ for Windows Version 3.0
| SEQ ID NO 5037
| LENGTH: 394
                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/960,253
CURRENT FILING DATE: 2001-09-20
NUMBER OF SEQ ID NOS: 187
SOFTWARE: FASCSEQ for Windows Version 4.0
SEQ ID NO 117
LENGTH: 398
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-960-253-117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Homo sapiens US-09-918-995-5037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-918-995-5037
                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 175
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Benson, Darin R.
APPLICANT: Mohamath, Raodoh
APPLICANT: Mohamath, Raodoh
APPLICANT: Lodes, Michael J.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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Local Similarity 100.0%; Pred. No. 1.3e-74;
les 175; Conservative 0; Mismatches 0;
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193
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                                                                                                                                                                                                                                                                           al Similarity
175; Conserv
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                               GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 2062
                                                                                                                                                                                                                  GTGTTTCTAGGTCGTGGCGTCGGGCTTCCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGC
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GAAATGCAGCGAGGATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG
                                                                                          GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAA
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                                                                                                                                                                                                                                                                         8.4%; Score 175; DB 9; 1 larity 100.0%; Pred. No. 1.3e-74; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                      Length 398;
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  247
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; ORGANISM: Homo sapiens
US-09-960-253-107
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; ORGANISM: Homo sapiens
US-09-833-790-349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 107, Application US/09960253
Patent No. US20020123619A1
GRMERAL INFORMATION:
APPLICANT: Benson, Darin R.
APPLICANT: Mohamath, Raodoh
APPLICANT: Mohamath, Raodoh
APPLICANT: Lodes, Michael J.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.512
CURRENT APPLICATION NUMBER: US/09/833,790
CURRENT FILING DATE: 2001-04-11
NUMBER OF SEQ ID NOS: 440
SOFTWARE: FBEUSEQ for Windows Version 4.0
SEQ ID NO 349
LENGTH: 521
TYPE: NAME
                                                                                                                                                                                                                                                                                                          PILE REFERENCE: 210121.556
CURRENT APPLICATION NUMBER: US/09/960,253
CURRENT FILING DATE: 2001-09-20
NUMBER OF SEQ ID NOS: 187
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 107
LENGTH: 665
                                                                                                                                       Query Match 8.4%; Score 175; DB 9; 1
Best Local Similarity 100.0%; Pred. No. 1.3e-74;
Matches 175; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%;
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APPLICANT: Wang, Tongtong
APPLICANT: Secrist, Heather
APPLICANT: Mohamath, Raodoh
APPLICANT: Indirias, Carol Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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1948 GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAA 2007
                                                                                            1888 GTGTTTCTAGGTCGTGGCGTCGGGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGC 1947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2008 GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 2062
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              52
                                             GTGTTTCTAGGTCGTGGCGTCGGGCTTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGC 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAA 2007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTGTTTCTAGGTCGTGGCGTCGGGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGC
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Pred. No. 1.3e-74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                            0; Indels
                                                                                                                                                                                         Length 665
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APPLICANT: Benson, Darin R.

APPLICANT: Mohamath, Raodoh

APPLICANT: Lodes, Michael J.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER

FILE REFERENCE: 210121.556

CURRENT APPLICATION NUMBER: US/09/960,253

CURRENT FILING DATE: 2001-09-20

NUMBER OF SEQ ID NOS: 187

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 106

LENGTH: 722
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; ORGANISM: Homo sapiens
US-09-960-253-106
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                                           US-10-084-817-316
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                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/084,817
CURRENT FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: 60/270,784
PRIOR FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 365
SOFTWARE: PERL Program
SEQ ID NO 316
TYPE-. NA 686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 316, 1
Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 106, Application US/09960253 Patent No. US20020123619A1
  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 8.4%; Score 175; DB 9; Best Local Similarity 100.0%; Pred. No. 1.3e-74;
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Susan Stuart
APPLICANT: Jed G. Nuchtern
APPLICANT: Sharon E. Plon
APPLICANT: Jason M. Shohet
TITLE OF INVENTION: GENES REGULATED BY MYCN ACTIVATION
FILE REFERENCE: PA-0046 US
                                                             FEATURE:
NAME/KEY: misc_feat
OTHER INFORMATION:
                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
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                                                               Incyte ID No. US20030119009A1 034181CB1
  8.4%;
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  Score 175;
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Length 3686;
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RESULT 13
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                                                                                                      Sequence 3, Application US/10097340 Publication No. US20030087250A1 GENERAL INFORMATION:
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US-09-864-864-300
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SOFTWARE: COTIXA Invention Disclosure Database SEQ ID NO 300
LENGTH: 3859
TYPE: DAA
ORGANITO:
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                                       APPLICANT: John MONAHAN
APPLICANT: Manjula GANN
APPLICANT: Sebastian HO
APPLICANT: Shubhangi KA
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APPLICANT:
APPLICANT:
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APPLICANT:
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APPLICANT: Behson, Darin R.
APPLICANT: Behson, Darin R.
APPLICANT: Catter, Darrick
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.523
CURRENT APPLICATION NUMBER: US/09/864,864
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                                                                                                                                                                                                                          GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 279
Steve G. KOVATS
Rachel E. MEYERS
                                  Sebastian HOERSCH
Shubhangi KAMATKAR
                                                                        Manjula GANNAVARAPU
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Secrist, Heather
Lodes, Michael J.
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RESULT 14
US-10-163-587A-3
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; ORGANISM: Homo sapiens
US-10-097-340-3
                                                                                                                                                                           Sequence 3, Application US/10163587A
Publication No. US20030096263A1
GENERAL INFORMATION:
APPLICANT: Oliveira, Marcos
TITLE OF INVENTION: SELECTIVE PARP-1 TARGETING
FILL REFERENCE: 50229-306
CURRENT APPLICATION NUMBER: US/10/163,587A
CURRENT FILING DATE: 2003-01-10
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PRIOR REPLICATION NUMBER: 05/276,025
PRIOR REPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR PELING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR APPLICATION NUMBER: 60/311,732
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/311,732
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
                                                                  SOFTWARE: PatentIn version 3.1 SEQ ID NO 3
                                                                                                              PRIOR APPLICATION NUMBER: 60/296,110 PRIOR FILING DATE: 2001-06-07 NUMBER OF SEQ ID NOS: 40
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PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
ORGANISM: Homo sapiens
                               TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 3859
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Karen GLATT
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Robert C. BAST, Jr.
Karen LU
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; Pred. No. 1.2e-74;
0; Mismatches 0;
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| Sequence 100, Application US/10334143 |
| Publication No. US20040009549A1 |
| GENERAL INFORMÁTION: | GOR VYACHESLAVOVICH |
| APPLICANT: GRIGGRIEV, IGOR VYACHESLAVOVICH |
| APPLICANT: SUDARSANAM, SUCHA |
| TITLE OF INVENTION: KINASES IDENTIFIED WITH THE METHOD |
| FILE REFERENCE: 038602/1543 |
| CURRENT APPLICATION NUMBER: US/10/334,143 |
| CURRENT FILING DATE: 2002-12-31 |
| PRIOR APPLICATION NUMBER: 60/343,169 |
| PRIOR FILING DATE: 2001-12-31 |
| VAMBER OF SEQ ID NOS: 207 |
| SOFTWARE: PATENTIN Ver. 2.1 |
| SEQ ID NO 100 |
| LENGTH: 3861 |
| TYPE: DNA |
| ORGANISM: Homo Bapiens |
Search completed: February 28, 2005, 10:48:47 Job time : 1121 secs
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; LOCATION: (160)..(
; OTHER INFORMATION:
US-10-163-587A-3
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                                                                                                                     2008 GAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG 2062
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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOPTWARR: FASSUSQ for Windows Version 4.0

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                                                TTTCCACCATCCAGAAGCAACTGGTGTTAATATCTGGAACATAAAATGGAATCACATCTT 37049
                                                                                                               AACACATTTTTTTTAATACAGAAAAGCTTAAAGGCACCAGAAGTGAAAAGTAACACACA 37109
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Pred. No. 4.7e-51;
0; Mismatches 575;
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                                  ACTGCACTCCAGCGCGGTGAGA
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                                                                                                                                                                                                                                                                                         GCCAGGTGCAGTGCCTGTAATCCCAGCACTTTGAGAGGCTGAGGCGGGGGGAT
                                                                   TGAGGCAGGAGAATCGCTTGAACACGGGAGGTGGAGATTGCAGTGAGCCAAGATCGCACC
                                                                                                                                                                           AAATTTTTTTTAATTAGCCAGTTGTGGTGAGCGCCTGTAGTCCCAGCTACTCGGGAGGC
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RESULT 3
US-09-949-016-12360
US-09-949-016-12360
Patent No. 6812339
Patent No. 6812339
GENERAL INFORMATION:
PATENTIAN:
ITITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND
FILL REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,756
PRIOR APPLICATION NUMBER: 60/241,768

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; ORGANISM: Human
US-09-949-016-12360
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PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 12360
LENGTH: 105679
Sequence 16409, Application US/09949016 Patent No. 6812339 GENERAL INFORMATION:
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Best Local S
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Pred. No. 7.2e-44;
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; ORGANISM: Human US-09-949-016-16409
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LENGTH: 107679
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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PRIOR FILING DATE: 2000-10-20
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Local Similarity 60.6%;
tes 452; Conservative
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AATTCCAGCACTTTGGGAGGTCGAGGTGGGCAGAT - - CATGAGGTCAGGAGTTCAAGACC
                            AATCCCAGCAATTTGGGAAGTCGAGGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACC
                                                                                       TAAATGCAACATCCTTTAGATAAAAATATTTGTCGGCCGGGCGCGGTGGCTCACGCCTGT
                                                                                                                               TGAAGTATAGTTCTATATTAAACAACAAGATCTAGGCCAGGTGCAGTGGCTCATGCCTGT
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Pred. No. 7.2e-44;
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806 AGCCTGGGCAACATGGAGAGATTCCC

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FILE REFERENCE: PXE-001

CURRENT APPLICATION NUMBER: US/09/792,616

CURRENT FILING DATE: 2001-02-23

NUMBER OF SEQ ID NOS: 27

SOFTWARE: PatentIn version 3.0

SEQ ID NO 1

LENGTH: 107820

TYPE: DNA

ORGANISM: Homo sapiens
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Best Local S
Matches 440
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APPLICANT: PXE International, Inc.
APPLICANT: University of Hawaii
TITLE OF INVENTION: Mutations in a gene encoding
TITLE OF INVENTION: Pseudoxanthoma Elasticum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application Patent No. 6780587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: "n" can be
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mes 440; Conservative
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AAACAACAAGATCTAGGCCAGGTGCAGTGGCTCATGCCTGTAATCCCCAGCAATTTTGGGAA
                                                CATATGCATGAGCATTCGAGTTTTTTTTGGAATATTCAGCAATTACAATCAAAAATGCCAT
                                                                                                              CTGGAATACAAGGTGAAGACACTGGGAAAAAGTCCTGGCCCCTAGCCCTAGCAAAAAACAAT
                                                                                                                                                                                                                                       CACGTTCTTCCTCTCTCTCTTCTTCCTAATGATGACATTTATGTACTGCACATGA
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                                                                              CATAGGATTAGAAAAGAAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATT
                                                                                                                                            AGCCCTGTCTGAGTGCCTTATGTTGTAATTGCTAAGCACCTGCCTATTGTTTTGCAAGGCC
                                                                                                                                                                                                        ACACTTTTTATGAGTTTTATTCATCTGCTTATTTTTCTTATCCTCTATACCAGAATTGAA
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Pred. No. 1.1e-43;
0; Mismatches 273;
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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR PILING DATE: 2000-10-3

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR PILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; NAME/KEY: misc_feature
; LOCATION: (1)...(36311)
; OTHER INFORMATION: n = A,T,C
US-09-949-016-13627
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US-09-949-016-13627/c
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SEQ ID NO 13627
LENGTH: 36311
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Best Local Similarity 54.8%;
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ORGANISM: Human
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version
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ACATCCTTGTCTTAAGCCACTAGCTTCTTCTCTATGGTTAACA---
                                              ATAATCACTCCAACCTCTTCCCTTTATTCCCCCAGCCCCAGGAATTGTAGCTGCTTCCTCC
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                                                                                            GITACTAACTTTTAGGATTCGGCCTATGTCTCACAACC----TTCTTGCTTACTCA 487
                                                                                                                                          AAGTGCTAGGATTACAAGCGTGAGCCACTGCGCCCCGGCCATTCTTCCAAGTTTCCTGGCT
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Pred. No. 4e-43;
0; Mismatches 550; Indels 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTGA----CATAATATGGTATCATTTAAGCATCAAAAAGAATGACTGCTATCTGCTGCTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGAGACCTCCCTGGGGAATACAGTGAG-----AACCTGTTCTCCACAAAAAGAAAAAG
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                                                                                                                                                                 GAGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGTGGGAGGATCGC-TGGGCTCAGGA
                                                                                                                                                                                                                                                                                                                                                                                      CACTTCGGGAGGCTGAGGCGTGAAGATCACTTGTAGCAGGAGTTTGAGACCAGTCTAGCC
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                                              GGACGCCTGTGGTCCCAGCTACTCAGGAGGCTGAAGTGGGAGAATGGCATGAACCCGGGA
                                                                                                                                                                                                                     AACTTGGCGAGACCCTGTCCCTAAAAAAATTTTTTTTTAATTAGCCAG----TTGTGGT
                                                                                                                                                                                                                                                                                                                                  AACTGTGGGAGGCTGAGGCAGGTGGATCACAAG-GTCAGGAGATCGAGACCATCCTGGCT
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    GGCCAGGCACAGTGGCTCATGCCTGTAATCCCAG

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RESULT

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i LENGTH: 38343
; TYPE: DNA
; ORGANISM: Human
; PEATURE:
; PEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(38343)
; OTHER INFORMATION: n = A
US-09-949-016-15714
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PATENT NO. 6812339

PATENT INFORMATION: Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN G.
TITLE OF INVENTION: WITH HUMAN DISEASE, MET
FILLE REFERENCES: CLOO1307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR PILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                            15839
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                                       177 CCAGTGGCACTATCATGGCTCACCACAGCCTCAACCTTCAGGGCTCAGGTGATCCTCCCA 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        117 TTCAAAATATCTTTTTTTTTTTTTTTGAGACAGGGTCACACTGTCACCCCAGGCTAGAGT
ACAACAAGATCTAGGCCAGGTGCAGTGGCTCATGCCTGTAATCCCAGCAATTTGGGAAGT 766
                                                                                                                       ACAACCTTCTTGCTTA-CTCAACATCCTTGTCTCTAAGCCACTAGCTTCTTCTCTATGG
                                                                                                                                                                                                                                                                                                                                                                          A--TGATCCACCCGCCTCAGCCTCCCAAAGTGCTAGGATTACAGGTGTGAGCCACCACAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAGGGATCCGGCCACCTCCAGCCTCCCAAAGTGCTAGGATTATAGGCATGAGCCACTGTGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TA------GAGACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTTCAGCCTCCCGAGTAGATGGGACTACAGGCACCTGCCACCACCCCCAGCTAATTTTTG
                                                                                TAGGATTAGAAAAGAAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATTAA
                                                                                                                                                                CCAG-CCTACCTTCAACGTATCTAACTGGTTACTAACTTTTAGGATTCGGCCTATGTCTC
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Pred. No. 4.6e-43;
0; Mismatches 357;
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Query Match Best Local Similarity 52.8%; pred. No. 1e-42; Matches 815; Conservative 0; Mismatches 641; Indels 88; Gaps 12; Oy 90 TCATATTTCCACTTATTCTAGTTTAGAAATATCTTTTTTTT	RESULT 8 US-09-949-016-12032 ; Sequence 12032, Application US/09949016 ; Sequence 12032, Application US/09949016 ; Patent No. 6812339 ; GENERAL INFORMATION: APPLICANT: VENTER, J. Craig et al. APPLICANT: VENTER, J. Craig et al. TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF ; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF ; CURRENT PILLICATION NUMBER: US/09/949,016 ; CURRENT PILLING DATE: 2000-04-14 ; PRIOR APPLICATION NUMBER: 60/241,755 ; PRIOR FILING DATE: 2000-10-20 ; PRIOR FILING DATE: 2000-10-03 ; PRIOR APPLICATION NUMBER: 60/237,768 ; PRIOR FILING DATE: 2000-10-03 ; PRIOR APPLICATION NUMBER: 60/231,498 ; PRIOR FILING DATE: 2000-09-08 ; NUMBER OF SEQ ID NOS: 207012 ; SOFTMARE: PastbEQ for Windows Version 4.0 ; SEQ ID NO 12032 ; LENGTH: 40655 ; TYPE: DNA ; ORGANISM: Human US-09-949-016-12032	Db 16139 NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
17 17 17 17 17 17 17 17 17 17 17 17 17 1	Qy 7 Db 166 Qy 7 Db 167 Qy 8 Db 167 Qy 9 Db 168 Qy 9 Db 168 Qy 9 Db 168	Qy 5 Db 164 Qy 6 Db 165 Db 165
	TECAGTGGCTCATGCCTGTAATCCCAGCAATTTGGGAAGTCGAGGTGGAAGTTGCTTG 786 647 TACAGTGGCTCACACCTGTAATCCCCAGCAATTTGGGAAGTCGAGGTGGAAGATCACTTTG 16706 647 TACAGTGGCTCACACCTGTAATCCCCAACACTTTGGGAAGACCAGCCAG	547 TATTCATCTGCTTATTTTTCTTATCCTATACCAGAATTGAATATTTTCAAATAAAGCA 606

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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

FITTLE OF INVENTEN: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF ITTLE OF INVENTEN: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 40655

TYPE: DNA

ORGANISM: Human

US-09-949-016-15919
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US-09-949-016-15919
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Best Local Sim.
Matches 815;
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                                                                                                                                                                                                                                                                                                                                  GTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCCGGCCACCTC
                                                                                                                                                                                                                                                                                                                                                                                         CCTGCCACCACCCCCAGCTAATTTTT------
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                                                                                                                              ТАТТСАТСТЭСТТАТТТТСТТАТССТСТАТАССАЗААТТЭААТАТТТТСАААТААЭССА
                                                                                                                                                         AGAATCCCAGCATCCTAACTCAGTCACTTATGATCTTCTGTATTTCCTTGGTCTTCCCTC
                                                                                                                                                                             AACATCCTTGTCTCTTAAGCCACTAGCTTCCTCTCTATGGTTAACACTTTTTATGAGTTT
                                                                                                                                                                                                           TTTTATTATCAAAACAATATTATGTTACGAAAA-----
                                                                                                                                                                                                                                   ATCTAACTGGTTACTAACTTTTAGGATTCGGCCTATGTCTCACAACCTTCTTGCTTACTC
                                                                                                                                                                                                                                                              GGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGTGCCCGGCTTA----GAACAT
                                                                                                                                                                                                                                                                                         AGCCTCCCAAAGTGCTAGGATTATAGGCATGAGCCACTGTGCCCAGCCTACCTTCAACGT
                                                                                                                                                                                                                                                                                                                   ACTTCCCAGGCTCAAGCAATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGATTACAGTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACCTTCAGGGCTCAGGTGATCCTCCCACTTCAGCCTCCCGAGTAGATGGGACTACAGGCA
 ATTTTTCCCTGTCACTCCCAGTGTTTCTGGTCATCATTTAAAACAATTATCTAGGCCAAG
                        ATTTTAATAAACTATATTTTGAAGTATAGTTCTATATTAAACAACAAGATCTAGGCCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 222.4; DB 4;
Pred. No. 1e-42;
0; Mismatches 641;
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US-09-949-016-14577
US-09-949-016-14577
Sequence 14577, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
TILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT APPLICATION NUMBER: 60/241,755

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; LOCATION: (1)...(678533
; OTHER INFORMATION: n =
US-09-949-016-14577
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Best Local
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ATCCGGTGAAACCTCGTCTACTAAAAAATACAAAAAAATTAGCCGAGCGTGGTGGCGGG
                                                           CTTTGGGAGGCCGAGGCGGAT--CACGAGGTCAGGAGATCGAGACCATCCTGGCTA
                                                                                       ATTTGGGAAGTCGAGGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCA 815
                                                                                                                           CCTTAACAGTTCTAGATÁCÁTTGÁGGCCGGGCACÁGTGGCTCAAGCCTGTÁATCCCAGCA
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                              CTTATCCCCTCTATAGCTCCCTGCTTGTGCAAGCTACCTTATCTCAGAATAGTTCCAGTT
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Pred. No. 1.5e-41;
0; Mismatches 662;
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PILE REFERENCE: CLOO1307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08
US-09-949-016-14578
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                                                                                                                                             SOFTWARE: FastSEQ for Windows Version SEQ ID NO 14578
                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION
         TYPE: DNA
ORGANIAM: Human
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(678533)
OTHER INFORMATION: n = A,
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Patent No. 681233
                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                              LENGTH:
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Matches 716; Conserv
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Pred. No. 1.5e-41;
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RESULT 12
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION
FILE REFERENCE: CL001107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ
SEQ ID NO 12513
                                                                                                                                                                                                                                                                                                               Matches 426;
                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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48329
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                                                                                                     CGAGTAGATGGGACTACAGGCACCTGCCACCCCCCAGCTAA-----TTTTTTGT
                                                                                                                                                                                          ATCATGGCTCACCACAGCCTCAACCTTCAGGGCTCAGGTGATCCTCCCACTTCAGCCTCC
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                                  AGAGACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCC
                                                                            TGAGTGGCTGGGACTACAAGCACACGCCACGCCTGGTTAATTTGTTGTATTTTAGT
                                                                                                                                                     ATCTAGGCTCACTGCAACCTCTGCCACCAGGGTTCACACAATTCTCCTGCCTCAGCCTCC
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                                                                                                                                                                                                                                                                                                             Score 219.2; DB 4;
Pred. No. 9.1e-42;
0; Mismatches 283;
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US-09-949-016-17050/c
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Window
SEQ ID NO 17050
LENGTH: 103988
                                                                                                                        Best Local Sim Matches 426;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17050, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                             Local Similarity
                                                       48509
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                                                                                                                      Score 219.2; DB 4;
Pred. No. 9.1e-42;
0; Mismatches 283;
                                                                                                                         Indels
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RESULT 14
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES AS:

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF

PILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR PILING DATE: 2000-10-03

PRIOR PILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012 SOFTWARE: FastSEQ for Windows Version SEQ ID NO 1286 FEATURE: misc_feature NAME/KEY: misc_feature LOCATION: (1)...(152582) TYPE: DNA
ORGANISM: Human LENGTH: 152582 ASSOCIATED
OF DETECTION AND USES THEREOF

Sequence 12086, Application US/09949016 Patent No. 6812339

GENERAL INFORMATION:

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RESULT 15
US-09-949-016-17390
Sequence 17390, App
Sequence 17390, App
Patent No. 6812339
GENERAL INFORMATION
APPLICANT: VENTER,
TITLE OF INVENTION
TITLE OF INVENTION
TITLE REFERENCE: CI
CURRENT APPLICATION
CURRENT APPLICATION
PRIOR APPLICATION
PRIOR FILING DATE:
PRIOR FILING DATE:
Sequence 17300, Application US/09949016
; Patent NO. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES AS:
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.
SEQ ID NO 17390
IENGTH: 152583
TYPE: DNA
ORGANISM: Human
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(152583)
OTHER INFORMATION: n = A,T,C or G
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Search completed: February 28, 2005, 01:15:34 Job time : 453.133 secs

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Run
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1: /cgn2_6/ptodata/2/pubpna/US07 PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

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5: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

6: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

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7: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

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11: /cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*

15: /cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*

16: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

16: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

17: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

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19: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

20: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

21: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

22: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

23: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

24: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

25: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

26: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

27: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*

28: /cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
      Match Length
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	e 176	899,	U	e 332	33297,	e 1776	e 70,	2768	34,	446,	Sequence 2326, Ap	238,	Sequence 661, App		۳,	177	e 100	06, A	Sequence 316, App	987	986,	987,	e 986,	Sequence 987, App	986,	e 12000,	e 11999	33300,		65	w -	Ü	00	Sequence 107, App

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CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 2085
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test
FILE REFERENCE: 18810-82152
                                                                                                                                                     Query Match 100.0%; Score 2085; Best Local Similarity 100.0%; Pred. No. 0; Matches 2085; Conservative 0; Mismatches
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ORGANISM: Homo sapiens
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                    TTCTCTAAGTAAAACTTTTATTTGTTCCATCATATTTTTCCACTTATTCTGTTTACCTTCA 120
TTCTCTAAGTAAAACTTTTATTTGTTCCATCATATTTTCCACTTATTCTGTTTACCTTCA
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	1141 TGCAGGGTCCACCTCGGGCCAATCAACTATATTCCCGAGGCGGGGGCCTGCGCTTCCCGG 1200
	1081 CGACGGTCTGTGGACGGCAGGTTAGAACGCCCGTCCAAGCCAGGAGGGTGGACCTAGCAC 1140
	1021 CTGGTATTTGTAGTAAATTCATAATAAAGGAAATGCTAGGTTTCAGTTGGTATTTTGTCC 1080
	961 CTAACATGAAAACGTCTGTGATGACTATTGCCCACAAAGTCACAGGTACTGCTAATACTC 1020
	901 CCACAATTTCGAGTAGTGATGAGCTTAATAATATTTCGAGTTATCACCAACAACTGTAAA 960
	841 CTTTACACACACACACACACACACACAAAATATCTGATAGCAACAGGTGCAGTCATTA 900
	781 TGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACATGGAGAGATTCCCCATCTCTTT 840
	721 GCCAGGTGCAGTGGCTCATGCCTGTAATCCCAGCAATTTGGGAAGTCGAGGTGGGAGGAT 780
	661 AAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATTAAACAACAAGATCTAG 720
	601 AAAGCACACTCATGTTACAATCTTTGAAATGGAAAAAAAA
	541 GAGTTTTATTCATCTGCTTATTTTTCTTATCCTCTATACCAGAATTGAATATTTTCAAAT 600
	481 TTACTCAACATCCTTGTCTCTTAAGCCACTAGCTTCTTCTCTATGGTTAACACTTTTTAT 540
	421 CAACGTATCTAACTGGTTACTAACTTTTAGGATTCGGCCTATGTCTCACAAACCTTCTTGC 480
	361 CACCTCAGCCTCCCAAAGTGCTAGGATTATAGGCATGAGCCACTGTGCCCAGCCTACCTT 420
	301 GACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCCGGC 360
	241 AGCCTCCCGAGTAGATGGGACTACAGGCACCTGCCACCCCCCCGGCTAATTTTTGTAGA 300
	181 TGGCACTATCATGGCTCACCACAGCCTCAACCTTCAGGGCTCAGGTGATCCTCCCACTTC 240
	121 AAATATCTTTTTTTTTTTTTTTTGAGACAGGGTCACACTGTCACCCAGGCTAGAGTCCAG 180

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Ş	1261	CATGCCCCTGATCCCAGCACTTCGGGAGGCTGAGGGCTGAAGATCACTTGTAGCAGGAGT	1320
B	1261	1 CATGCCCCTGATCCCAGCACTTCGGGAGGCTGAAGGCGTGAAGATCACTTGTAGCAGGAGT	1320
Ş	1321	TTGAGACCAGTCTAGCCAACTTGGCGAGACCCTGTCCCTAAAAAAAA	1380
뮰	1321	TIGAGACCAGICTAGCCAACTIGGCGAGACCCTGTCCCTAAAAAAATTTTTTTTTAATT	1380
Ş	1381	AGCCAGTTGTGGTGAGCGCCTGTAGTCCCCAGCTACTCGGGAGGCTGAGGTGGGAGGATCG	1440
Db	1381	AGCCAGTTGTGGTGAGCCCTGTAGTCCCCAGCTACTCGGGAGGCTGAGGTGGGAGGATCG	1440
Ş	1441	TGGGCTCAGGAGTTCCAGACTGCAGTGAGCCATGATGGCGGCACTGCACTCCAGCGCGG	1500
Db	1441	CTGGGCTCAGGAGTTCCAGACTGCAGTGAGCCATGATGGCGGCACTGCACTCCAGCGCGG	1500
γQ	1501	TGAG	1560
DЬ	1501	TGAGACTCAGTCTCAAAAATAAAAGGGGGAGGGGTTGGGGGTAAAATTAGTTGTGAAAT	1560
8	1561	AAG	1620
ДЪ	1561	. AAGTAAGACTTCCTGGGACAGAACAATCAAAGGGGTGGCGCCGGGTCCTCCAAAGAG	1620
γQ	1621	TAGCTCAGCCCAAGCCCCGCCTCGGCCCCAGGGCAGGCCGCCAGAGCTCCACCCGGC	1680
Db	1621	. CTAGCTCAGCCCAAGCCCCGCCTCGGCCCCCAGGGCAGCGGCCGCAGAGCTCCACCCGG	1680
γQ	1681	- AGC	1740
В	1681	ageceleccedeanactecececcecedecedecadecedecedecedecedecede	1740
Ş	1741	1 TGGACGCGGGTTCCCGTGGCGTTCCCGCGGCCATCAGCAATCTATCAGGGAACGGCG	1800
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Ş	1801	1 GTGGCCGGTGCGGCGTTCGGTGCGCTCTGGCCGCTCAGCCGTGGCGGCGGCTGAGCG	1860
Db	1801	. GTGGCCGGTGCGGCGTTCTGGCCGCTCAGCCGTTGGCGGCTGGGTGAG	186
Ş	1861		1920
Db	1861	CACGCGAGGCGAGGCGCAAGCGTGTGTTTCTAGGTCGTGGCGTCGGGCTTCC	1920
Ş	1921	1 CTTTGGCGGCAGCTAGGGGAGATGGCGGAGTCTTCGGATAAGCTCTATCGAGTCGAGTA	1980
qa	1921	1 CTTTGGCGGCAGCTAGGGGAGGATGGCGGAGTCTTCGGATAAGCTCTATCGAGTCGAGTA	1980
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뮹		1 GCCAAGAGCGGGCGCCTCTTGCAAGAAATGCAGCGAGGAGCATCCCCAAGAGTC	

US-10-239-676-1
US-10-239-676-1
Sequence 1, Application US/10239676
Sequence 1, Application WS/10239676
Publication No. US20030082609A1
GENERAL INFORMATION:
APPLICANT: DIEENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with Gene Regulation
FILE REFERENCE: 5013.1003
CURRENT APPLICATION NUMBER: US/10/239,676
CURRENT FILING DATE: 2002-09-24
PRIOR APPLICATION NUMBER: PCT/EP01/03968

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PRIOR FILING DATE: 2001-04-06

2000-04-06

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2000-09-01

NUMBER OF SEQ ID NOS: 228

SEQ ID NO 1

LENGTH: 10619

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: chemically tr

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77.2%;
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                                           CCGCCCCGTGGACGCGGGTTCCGT-GGCGTTCCCGCGGGCCAGGCATCAGCAATCTATCAG
                                                                ATTCGGTAGGCGTTCGGAAATTTCGTTTTTCGGTCGGTAGGGGGCGCGCGTCGTCGGTT
                                                                       GGTAGTATATGTTTTTGATTTTAGTATTTCGGGAGGTTGAGGCGTGAAGATTATTTGTAG
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QY 301 GACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCCGGC 360	3294 TGTATTATTATGTTTATATTTTAATTTTTAGGTTTAAGGTTAAGTTTTTT	Db 3236 AAATATTTTTTTTTTTTTTTTAGATARGGTTATATTTTTTTTTT	Db 3176 TTTTTTAAGTAAAATTTTTTTTTTTATTATTTTTTTTTT	Db 3116 TTTÄGGGÄTGÄTÄTÄĞTTÄÄTTTÄĞÄGÄTGĞTATGATTATGTTTTTTGATTTTGGTTA 3175 OY 61 TTCTCTAAGTAAAACTTTTATTTGTTCCATCATATTTTCCACTTATTCTGTTTACCTTCA 120	; Conservative 0; Mismatches TTTAGGGATGATATAGTTGTCAACCCAGAGATGG	Query Match 58.7%; Score 1223.6; DB 15; Length 10619; Best Local Similarity 77.2%; Pred. No. 4.2e-289;	; ILES: DAM ; ORGANISM: Artificial Sequence ; FEATURE: FEATURE: OFFIER INFORMATION: Chemically treated genomic DNA (Homo sapiens) [16-10-311-465-43]	8 2 7	PRIOR PRIOR PRIOR	CURREN CURREN PRIOR	; APPLICANT: PIEPENBROCK, Christian ; APPLICANT: BERLIN, Kurt ; TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Determ ; TITLE OF INVENTION: Cytosine methylation	; Sequence 43, Application US/10311455 ; Publication No. US20030143606A1 ; GENERAL INFORMATION: ; APPLICANT: OLEK, Alexander	RESULT 3 US-10-311-455-43	OY 2030 AAGGACTCGCTCCGGATGGCCATCATGGTGCAGGTGCGGGCCGCTGTGCGGGGGGG 2085	QY 1970 CGAGTCGAGTACGCCAAGAGCGGGCGCCTCTTGCAAGAAATGCAGCGAGAGCATCCCC 2029	OY 1910 GGCTTCCGGAGCTTTGGCGCAGCTAGGGGAGGAGGAGGAGTCTTCGGATAAGCTCTAT 1969	Qy 1850 CTGGGTGAGCGCACGCGAGGCGGCAGGCGCAAGCGTGTGTTTCTAGGTCGTGGCGTCG 1909
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APPLICANT: DIEK, ALEXANDER
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
ITTLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Transcription
ITTLE OF INVENTION: With DNA TRANScription
ITTLE OF INVENTION: With DNA TRANScription
ITTLE OF INVENTION: With DNA TRANScription
ITLE REFERENCE: 5013.1009
CURRENT EILING DATE: 2002-10-02
PRIOR APPLICATION NUMBER: US/10/240,453
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 350
LENGTH: 10619
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; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
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Sequence 1, Application US/10240589C

| Publication No. US20040076956A1
| GENERAL INFORMATION: |
| APPLICANT: OLEK, Alexander |
| APPLICANT: PIEPENBROCK, Christian |
| APPLICANT: BERLIN, Kurt |
| TITLE OF INVENTION: Diagnosis of Diseases Associated wit |
| TITLE OF INVENTION: Diagnosis of Diseases Associated wit |
| TITLE OF INVENTION: DIAGNOSIS of DISEASES ASSOCIATED NUMBER: US/10/240,589C |
| CURRENT APPLICATION NUMBER: US/10/240,589C |
| CURRENT APPLICATION NUMBER: PCT/EPO1/03972 |
| PRIOR APPLICATION NUMBER: DE 10019058.8 |
| PRIOR APPLICATION NUMBER: DE 10019058.8 |
| PRIOR APPLICATION NUMBER: DE 10019173.8 |
| PRIOR APPLICATION NUMBER: DE 10019173.8 |
| PRIOR APPLICATION NUMBER: DE 10032529.7 |
| PRIOR APPLICAT
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SEQ ID NO 1
LENGTH: 10619
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Best Local
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: chemically
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Pred. No. 4.2e-289;
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APPLICANT: DIEKNERCCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Ass
FILE REFERENCE: S013.1003
CURRENT APPLICATION NUMBER: US/10/239,676
CURRENT FILING DATE: 2002-09-24
PRIOR APPLICATION NUMBER: PCT/EP01/03968
DE 10019058.8
DE 10019058.8
DE 10019058.8
DE 10043826.1
PRIOR FILING DATE: 2001-04-06
2000-04-06
2000-04-07
2000-06-30
2000-09-01
NUMBER OF SEQ ID NOS: 228
SEQ ID NO 2
LENGTH: 10619
TYPE: DNA
OFGANTOM: Artificial Sections
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Sequence 2, Application US/10239676
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RESULT 7
US-10-311-455-44/c
; Sequence 44, Application US/10311455
; Publication No. US20030143606A1
; GENERAL INFORMATION:
; GENERAL INFORMATION:
; APPLICANT: DIEFENBROCK, Christian
; APPLICANT: BERLIN, Kurt

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; TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune Syst TITLE OF INVENTION: cytosine methylation FILE REFERENCE: 5013.1014
; CURRENT APPLICATION NUMBER: US/10/311,455
; CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: US/10/7537
; PRIOR APPLICATION NUMBER: DET/EP01/07537
; PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10032529.7
; PRIOR APPLICATION NUMBER: DE 10043826.1
; PRIOR APPLICATION NUMBER: DE 10043826.1
; PRIOR APPLICATION NUMBER: DE 10043826.1
; NUMBER OF SEQ ID NOS: 2424
; SEQ ID NO 44
; LENGTH: 10619
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; ORGANISM: Artificial Sequence;
; FEATURE:
; OTHER INFORMATION: chemically treated
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Pred. No. 4.3e-284;
0; Mismatches 463; I
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APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BEELIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Dy Means of Assessing the Methylation Status of (
TITLE OF INVENTION: With DNA TRANSCRIPTION
FILE REFERENCE: 5013.1009
CURRENT APPLICATION NUMBER: US/10/240,453
CURRENT FILING DATE: 2002-10-02
PRIOR APPLICATION NUMBER: PCT/EP01/03973
PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 1001913.8
PRIOR TILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR PILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR PILING DATE: 2000-09-01
PRIOR APPLICATION NUMBER: DE 10043826.1
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PRIOR APPLICATION NUMBER: DE 10043826.1
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US-10-240-453-2/c
19-10-240-453-2/c
; Sequence 2, Application US/10240453
; Publication No. US20030148326A1
; GENERAL INFORMATION:
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; SEQ ID NO 2
; LENGTH: 10619
; TYPE: DNA
; ORGANUISM: Artificial Sequence
; PEATURE: INFORMATION: chemically treated genomic DNA (Homo US-10-240-453-2
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Pred. No. 4.3e-284;
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Sequence 2, Application US/10240589C

Publication No. US20040076956A1

GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated wit
TITLE OF INVENTION: DNA repair
FILTE REFERENCE: 5013.1008

CURRENT APPLICATION NUMBER: US/10/240,589C

CURRENT FILING DATE: 2003-09-02

PRIOR APPLICATION NUMBER: PCT/EP01/03972

PRIOR APPLICATION NUMBER: DE 10019058.8

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR FILING DATE: 2000-04-06

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR FILING DATE: 2000-04-06

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR FILING DATE: 2000-04-07

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR FILING DATE: 2000-06-30
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; OTHER INFORMATION:
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PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 148
SEQ ID NO 2
LENGTH: 10619
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 1600; Conser
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                                                                  TGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACATGGAGAGATTCCCCCATCTCTTT
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Pred. No. 4.3e-284;
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	6667 CTTTACACACACACACACACACACACACACACAAAATATTCTAATAA
RESULT 10 US-10-027-632-154183 ; Sequence 154183, Application No. US20020198 ; GENERAL INFORMATION: Ident APPLICANT: Wang, David G. TITLE OF INVENTION: Ident FILTE OF INVENTION: Ident FILTE OF INVENTION: Poly FILE REFERENCE: 108827.12 CURRENT APPLICATION NUMBER: CURRENT APPLICATION NUMBER: PRIOR FILLING DATE: 2000-0. PRIOR APPLICATION NUMBER: PRIOR FILLING DATE: 2000-0. PRIOR APPLICATION NUMBER: PRIOR FILLING DATE: 1099-0.0 PRIOR APPLICATION NUMBER: PRIOR FILLING DATE: 1999-11 PRIOR APPLICATION NUMBER: PRIOR FILLING DATE: 1999-11 PRIOR APPLICATION NUMBER: PRIOR FILLING DATE: 1999-12 PRIOR APPLICATION NUMBER: PRIOR FILLING DATE: 1999-13 SERGITH: 844 TYPE: DNA ORGANIEM: Human US-10-027-632-154183 Query Match SEQ ID NO 154183 QUERY MATCH DE 1207 CTGCCCTCAGGGGAG QY 1207 CTGCCCTCAGGGGAG QY 1267 CCTGATCCCAGGCACT QY 1387 TTGTGGTGAGCGCCTC QY 1387 TTGTGGTGAAGATTAA AND DD 1507 TCAGTCTCAAAAATAA DD 301 TCAGTCTCAAAAATAA ON 1507 GACTTCCTGAGAATTAA ON 1507 GACTTCCTGAGAATTAA ON 1507 GACTTCCTGAGAAATAAA ON 1507 GACTTCCTGAGAAATAAA ON 1507 GACTTCCTGAGAAATAAA ON 1507 GACTTCCTGAGAAATAAAATAA ON 1507 GACTTCCTGAGAAATAAAATAA ON 1507 GACTTCCTGAGAAATAAAATAAA ON 1507 GACTTCCTGAAAATAAAATAAA ON 1507 GACTTCCTGAGAAAATAAAATAAA ON 1507 GACTTCCTGAGAAATAAAATAAA ON 1507 GACTTCCTGAAAATAAAATAAA ON 1507 GACTTCCTGAAAATAAAATAAA ON 1507 GACTTCCTGAAAATAAATAAAATAAAATAAAATAAA	Qy 1970 CGAG: Db 5532 CGAA: Qy 2030 AAGGJ
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APPLICANT: Waid, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
FILE REFERENCE: 108627.129

CURRENT APPLICATION NUMBER: US/10/027,632

CURRENT APPLICATION NUMBER: US 60/218,006

PRIOR APPLICATION NUMBER: US 60/218,006

PRIOR FILING DATE: 2000-07-12

PRIOR APPLICATION NUMBER: US 60/198,676

PRIOR PRILING DATE: 2000-04-20

PRIOR APPLICATION NUMBER: US 60/193,483

PRIOR FILING DATE: 2000-03-29

PRIOR APPLICATION NUMBER: US 60/185,218

PRIOR FILING DATE: 2000-03-29

PRIOR APPLICATION NUMBER: US 60/185,218

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-11-23

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR FILING DATE: 1999-11-23

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR FILING DATE: 1999-08-09

PRIOR FILING DATE: 1999-08-09

PRIOR FILING DATE: 1999-08-09
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US-10-027-632-154183
                                                                                                                           ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-154183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 154183, Application US/10027632
Publication No. US20030204075A9
GENERAL INFORMATION:
                                       Query Match
Best Local Similarity
Matches 822; Conserv
                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 154183
LENGTH: 844
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                                         Score 699.4; DB 17;
Pred. No. 4.9e-161;
1; Mismatches 12;
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GENERAL INFORMATION:

APPLICANT: Benson, Darin R.

APPLICANT: Mohamath, Raodoh

APPLICANT: Mohamath, Raodoh

APPLICANT: Lodes, Michael J.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER

FILE REFERENCE: 210121.556

CURRENT APPLICATION NUMBER: US/09/960,253

CURRENT FILING DATE: 2001-09-20
                                                                                                                                                                               Sequence 107, Application US/09960253
Patent No. US20020123619A1
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APPLICANT: Mannion, Jane
APPLICANT: Mannion, Jane
APPLICANT: Benson, Darin R.
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.523
CURRENT APPLICATION NUMBER: US/09/864,864
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 341
SOFTWARE: Corixa Invention Disclosure Database
SEQ ID NO 300
LENGTH: 3859
TYPE: DNA
TYPE: DNA
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; ORGANISM: Homo sapiens
US-09-960-253-107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 187
SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 107
LENGTH: 665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 12.7
Best Local Similarity 97.2
Matches 281; Conservative
                                                                                                                                                      Query Match
Best Local S
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                           ORGANISM: Homo sapiens -09-864-864-300
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APPLICANT:
APPLICANT:
                                                                                                                                    y Match 12.5%; Local Similarity 97.2%; hes 277; Conservative
                                                                                         1782 AATCTATCAGGGAACGGCGGTGGCCGGTGCGGCGTTTCGGTGCGCTCTGGCCGCTCAGC 1841
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Secrist, Heather
Lodes, Michael J.
Algate, Paul A.
Fling, Steve P.
                       GAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAGGTGC 2066
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 CCGTGCGGCTGGGTGAGCGCACGCGAGGCGGCGAGGCGCAAGC--GTGTTTCTAGGTCG
                                                                     AATCTATCAGGGAACGGCGGTGGCCGGTGCGGCGCTTCGGCCGCTCAGG
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Pred. No. 1.4e-54;
0; Mismatches 6
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                                                                                                                                    Score 261.4; DB 9; Pred. No. 3.4e-53; 0; Mismatches 6;
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TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REPERENCE: MRI-030
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-14
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR FILING DATE: 2001-09-14
PRIOR APPLICATION NUMBER: 60/314,967
PRIOR APPLICATION NUMBER: 60/311,732
PRIOR APPLICATION NUMBER: 60/311,732
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR APPLICATION NUMBER: 60/315,102
PRIOR APPLICATION NUMBER: 60/325,102
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APPLICANT: John MONAHAN
APPLICANT: Manjula GANNA
APPLICANT: Sebastian HOE
APPLICANT: Shabhangi KAM
APPLICANT: Steve G. KOVA
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US-10-097-340-3
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-3
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SEQ ID NO 3
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                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 277; Conserv
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                            CGTGGCGGCTGGGTGAGCGCACGCGAGGCGGCGAGGCGGCAAGCGTGTGTTTCTAGGTCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAGGTGC 2066
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CCGTGCGGCTGGGTGAGCGCACGCGAGGCGAGGCGAAGC--GTGTTTCTAGGTCG
                                                                                              AATCTATCAGGGAACGGCGGTGCCGGTGCGGCGTGTTCGGTGCGCTCTGGCCGCTCAGG
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Karen GLATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peter VEIBY
Gordon B. MILLS
Robert C. BAST, Jr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sebastian HOERSCH
Shubhangi KAMATKAR
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                                                                                                                                                                                          Score 261.4; DB 14; Pred. No. 3.4e-53; 0; Mismatches 6;
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PRIOR APPLICATION NUMBER: 60/296,110
PRIOR FILING DATE: 2001-06-07
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
LENGTH: 3859
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (160)...(3204)
OTHER INFORMATION:
US-10-163-587A-3
Search completed: February 28, 2005, 06:52:37 Job time : 5156.4 secs
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US-10-163-587A-3
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Best Local Similarity
Matches 277; Conserv
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TITLE OF INVENTION: SELECTIVE PARP-1 TARGETING FOR DESIGNING CHEMO/RADIO SENSITIZING
FILE REFERENCE: 50229-306
CURRENT APPLICATION NUMBER: US/10/163,587A
CURRENT FILING DATE: 2003-01-10
CURRENT FILING DATE: 2003-01-10
                                                                                                                                                                                1962 AGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCCCCTCTTGCAAGAAATGCAGCGAGA 2021
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ilarity 97.2%;
Conservative
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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Query
Match Length
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
      US-09-909-317-7
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        GenCore version 5.1.6 (c) 1993 - 2005 Compugen
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                     US-08-222-177A-140
US-08-222-177A-397
US-08-222-177A-137
US-08-222-177A-13
US-08-222-177A-13
US-08-222-177A-403
US-08-222-177A-43
US-08-222-177A-23
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Sequence 140, App Sequence 198, App Sequence 197, App Sequence 117, App Sequence 400, App Sequence 403, App Sequence 403, App Sequence 232, App Sequence 232, App Sequence 232, App Sequence 241, App Sequence 241, App Sequence 211, App Sequence 101, App Sequence 101, App Sequence 101, App Sequence 226, App Sequence 211, App Sequence 31, App Sequence 221, App Sequence 31, App Sequence 361, App Sequence 351, App Sequenc
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US-08-222-177A-140

Query Match 100.0%; Score 36; DB 1; Length 37; Best Local Similarity 100.0%; Pred. No. 7.8e-05; Matches 36; Conservative 0; Mismatches 0; Indels

0

Gaps

US-08-222-177A-140 US-08-222-177A-140 Sequence 140, Application US/0 Patent No. 5582979 GENERAL INFORMATION: INFORMATION OF SEQUENCES: INFORMATION INFORMATION: INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS: INFORMATION: INFORMATION INFORMATION: INFORMATION INFORMATION: INFORMATION: INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS: INFORMATION: INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS: INFORMATION: INFORMATION: INFORMATION INFORMATION: INFOR	28 36 100.0 5 29 36 100.0 5 30 36 100.0 5 31 36 100.0 5 32 36 100.0 5 33 36 100.0 5 34 36 100.0 5 36 100.0 5 37 36 100.0 5 38 36 100.0 5 41 36 100.0 5 42 36 100.0 5 43 36 100.0 5 41 36 100.0 5 42 36 100.0 5 43 36 100.0 5 44 1 36 100.0 5 45 36 100.0 6
Jation US/08222177A N: James L. N: LENGTH POLYMORPHISMS IN N: (AC-dA)n.(dG-dT)n SEQUENCES AND DES: 460 DES: 460 DES: 460 DES: 460 DES: 460 DES: 541tt Ross & Stevens, S.C. Sxcelsior Drive, Suite 401 Sxcelsior Drive, Suite 401 Sin 14 15 FORM: 10 FORM: 10 FORM: 10 SM: PC-DOS/MS-DOS SMITIN Release #1.0, Version #1.25 INDATA: 435 VERTION: MBER: US 07/341,562 US 07/341,562 US 08/222,177A 144: MBER: 150 MBER: 10 MBER:	50 1 US-08-222-177A-186 50 1 US-08-222-177A-328 50 1 US-08-222-177A-367 50 1 US-08-222-177A-379 51 1 US-08-222-177A-317 51 1 US-08-222-177A-325 51 1 US-08-222-177A-337 51 1 US-08-222-177A-391 53 1 US-08-222-177A-304 55 1 US-08-222-177A-304 56 1 US-08-222-177A-394 57 1 US-08-222-177A-394 58 1 US-08-222-177A-394 59 1 US-08-222-177A-394 60 1 US-08-222-177A-394 61 US-08-222-177A-394 62 6 5506118-3 62 6 5506118-3
METHODS OF USING SAME	Sequence 186, App Sequence 328, App Sequence 367, App Sequence 379, App Sequence 317, App Sequence 301, App Sequence 307, App Sequence 307, App Sequence 318, App Sequence 318, App Sequence 324, App Sequence 324, App Sequence 324, App Sequence 344, App Sequence 344

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CURKENT COURTON APPLICATION APPLICATION 435
FILING DATE:
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
PILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.60
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 10080831-2100
TT.REPAN: (608) 831-2106
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US-08-222-177A-198
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                                    Sequence 397, Application US/08222177A
PATENT NO. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PATENT NO. 5582979
PATENT NO. 558297
PATEN
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Best Local Similarity
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TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genom
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING BYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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            ADDRESSEE:
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CITY: Madison
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DeWitt Ross & Stevens, S.C.
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                                                                                                                                                                                                                                                                                                                                                          Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence 137, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOPTIMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
FILING DATE:
CLASSIFICATION 235
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 09865.601
TELECOMUNICATION TORNEYION:
TELEPHONE: (608) 831-2106
TELEPTAX: (608) 831-2106
                                                                                                                       ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPUTER:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, VE
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 38 base pairs
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILLING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                       FILING DATE:
                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                  CITY: Madison
STATE: Wiscons
                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 100.0%;
Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
CITY: M
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                                                                                                      US/08/222,177A
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Pred. No. 7.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                           Version #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37
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                                                                                                                                                                                                                                                                                                                                                                     AND METHODS
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RESULT 5
US-08-222-177A-119
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                            FILING DATE:

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/341,562

PILING DATE: 21-APR-1989

ATTORNEY/AGENT INFORMATION:

NAME: Sara, Charles S.

REGISTRATION NUMBER: 30,492

REFERENCE/DOCKET NUMBER: 30,492

REFERENCE/DOCKET NUMBER: 09865.601

TELECOMMUNICATION INFORMATION:

TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
CLONE: mfd29rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
IMMEDIATE SOURCE:
CLONE: mfd23rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 39 base pairs
                                    TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 53717-1914
                                                                      STRANDEDNESS:
                                                                                     TYPE: nucleic acid
                                                                                                                                                                                   TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE! DeWitt Ross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity
36; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BOOO Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Weber, James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                     linear
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                                    DNA (genomic)
                                                                          double
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US-08-222-177A-403
; Sequence 403, Application US/08222177A
; Patent No. 5582979
; GENERAL INFORMATION: James L.
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPH;
TITLE OF INVENTION: (dC-dA)n.(dG-dT);
NUMBER OF SEQUENCES: 460
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US-08-222-177A-400
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                                                                                                                                                            RESULT 7
                                                                                                                                                                                                                                                                                                                                                       ; MOLECULE TYPE: US-08-222-177A-400
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Matches
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                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Weber, James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: LENGTH POLYMORPHISMS IN TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 40 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Madison
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                                                                                                                                                                                                                                                                                    36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36;
                                                                                                                                                                                                                                     1 CACACACACACACACACACACACACACACACACACA 36
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8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                                                                                                                                                                                                                                                    100.0%; Score 36; DB 1; ilarity 100.0%; Pred. No. 7.8e-05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
 ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                       DNA (genomic)
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                                 LENGTH POLYMORPHISMS IN (dC-dA)n, (dG-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07/341,562
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Version
                                                                                                                                                                                                                37
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                                                                                                                                                                                                                                                                                                                     Length 40;
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                                                                                                                                                                                                                                                                                     Indels
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ADDRESSEE:

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US-08-222-177A-74
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                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 74, Application US/08222177A Patent No. 5582979 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -08-222-177A-403
                                                                                                     ZIP: 53717-1914

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
TOPOLOGY: lir
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                               CITY: Madison
                                                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
36; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CACACACACACACACACACACACACACACACACA 36
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                                                                                                                                                                                                                                              Wisconsin
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                                                                                                                                                                                                                                                                           3: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                 Weber, James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (608) 831-2106
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E: DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                           (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME
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                                                                                                                                                                                                                                                                                                                              460
                                                                                           US/08/222,177A
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US-08-222-177A-183
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Patent NO. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMOR:
TITLE OF INVENTION: (dC-dA)n.(dG-d-dT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS LENGTH: 41 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
              IMMEDIATE SOURCE: CLONE: mfd45rs
                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 53717-1914

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOSTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                  MOLECULE TYPE:
                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
TELEPHNE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: 1
MOLECULE TYPE:
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CLONE: mfd8rs
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                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                 STRANDEDNESS:
                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                    TELEFAX:
                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Madison
STATE: Wiscon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
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8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
                                                                                                                                                                                    (608) 831-2106
                                                               linear
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                                                                                                                                                    183:
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                                                                                                                                                                                                                                       09865.601
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RESULT 10
US-08-222-177A-232
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                                                                                     RESULT 11
US-08-222-177A-53
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                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Sequence 53, Application US/08222177A
Patent No. 5582379
PATENT INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 5582979
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 53717-1914

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: IBM FC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DI
IMMEDIATE SOURCE:
CLONE: mfd63rs
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 41 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
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8000 Excelsior Drive, Suite 401
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                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                    DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                         double
                                                                                                                                                                                                                                              100.0%; Score 36; DB 1; 100.0%; Pred. No. 7.8e-05;
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US-08-222-177A-229
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                                                                                                                                                                                                                                                                                                                Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                   Sequence 229, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
PILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SETA, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.
CURRENT APPLICATION NUMBER: US/08/222,177A
APPLICATION NUMBER: US/08/222,177A
            COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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IMMEDIATE SOURCE:
CLONE: mfdlrs
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ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                           TITLE OF INVENTION: LE
TITLE OF INVENTION: (d
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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STREET: 0...
Madison
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                                                                                                                                                                             STATE:
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ZIP: 53717-1914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX:
FILING DATE:
                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wisconsin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           42 base pairs
                                                                                                                                                                                                          B: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (608) 831-2106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double
                                                                                                                                                                                                                                                                                                                    James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 36; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                              LENGTH POLYMORPHISMS IN (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:

US 07/341,562

CLASSIFICATION:

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RESULT 13
US-08-222-177A-388
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY,AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 09.492
REFERENCE/DOCKET NUMBER: 09.65.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION: James L.

APPLICANT: Weber, James L.

TITLE OF INVENTION: LENGTH POLYMORPHISMS IN

TITLE OF INVENTION: (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING
                                                                  TELEX:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                        ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                  SEQUENCE CHARACTERISTICS:
LENGTH: 43 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SATA, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION: 435
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STRANDEDNESS:
                                                                                                              TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLONE: mfd62rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (608) 831-2106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 36; DB 1; 100.0%; Pred. No. 7.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     460
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RESULT 15 US-08-222-177A-241

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; TOPOLOGY: linear
; MOLECULE TYPE: DNA
; IMMEDIATE SOURCE:
; CLONE: mfd49rs
US-08-222-177A-195
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                                                                                Query Match 100.0%;
Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 195, Application US/08222177A Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
                                                                     Matches
                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/222,177A FILING DATE: CLASSIFICATION: 435
CLASSIFICATION DATA: APPLICATION DATA: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/ACENT INFORMATION: NAME: Sara, Charles S. REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: DNA IMMEDIATE SOURCE: OCLONE: mfd123rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: LENGTH POLYMORPHISMS IN TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                TELEX:
                                                                                                                                                                                                                          STRANDEDNESS: double
                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                      TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
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                       53717-1914
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CACACACACACACACACACACACACACACACACA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                     Conservative
                                                                                                                                                                                                           linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US/08/222,177A
                                                                   0
                                                                                  Score 36; DB 1; 1
Pred. No. 7.8e-05;
                                                                   Mismatches
                                                                                                   Length 44;
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                                                                     Indels
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TELEX:
INFORMATION FOR SEQ ID NO: 24
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genom
IMMEDIATE SOURCE;
CLONE: mfd66rs
US-08-222-177A-241
Search completed: February 28, 2005, 01:15:34 Job time: 7.32316 secs
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION UMBER: US/08/222,177A
FILING DATE:
CLASSIFICATION UMBER: US/08/222,177A
FILING DATE:
CLASSIFICATION UMBER: US 07/341,562
FILING DATE: US 07/341,562
FILING DATE: 1-APPLICATION:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGÉNT INFORMATION:
NAME: SATA, Charles S.
REGISTRATION UMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELEPHONE: (608) 831-2106
TELEPHONE: (608) 831-2106
TELEPAX: (608) 831-2106
                                                                                                                                                                                Query Match
Best Local Similarity
Matches 36; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 241, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Weber, James L.
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESS: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
CITY: Madison
STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 53717-1914
                                                                                                                                                                              ch 100.0%; Score 36; DB 1; Length 44; l Similarity 100.0%; Pred. No. 7.8e-05; 36; Conservative 0; Mismatches 0; Indels
                                                                                                                     1 CACACACACACACACACACACACACACACACACA 36
                                                                                        CACACACACACACACACACACACACACACACACACA 37
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Run
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Published Applications NA:*

1: /cgn2_6/ptodata/2/pubpna/USO7_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/USO6_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/USO6_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/USO7_NEW_PUB.seq:*

6: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*

1: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*

7: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*

1: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IDENTITY_NUC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             February 28, 2005, 00:47:38; Search time 88.8932 Seconds (without alignments) 2399.695 Million cell updates/sec
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36
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Copyright (c) 1993 - 2005 Compugen Ltd.
/ Cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq: *
/ Cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq: *
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6/ptodata/2/pubpna/US09A
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

0 0	Result No.
	Score
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US-09-852-903C-23 US-09-909-317-7 US-09-852-903C-24 US-09-852-903C-25 US-09-852-903C-25 US-10-661-088-24 US-10-661-097-24 US-10-661-355-24 US-10-661-415-24 US-10-661-415-24 US-10-661-415-24 US-10-661-415-24	SUMMARIES
Sequence 23, Appl Sequence 7, Appli Sequence 74, Appl Sequence 678, App Sequence 678, Appl Sequence 24, Appl	Description

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Sequence 26864, A Sequence 7594, Ap Sequence 53, Appl	e 37, App e 450, App e 38, App e 2662, A e 5212, A e 6881, A	35, 34, 12, 56, 56, 56, 56, 56, 56, 56, 56, 56, 56	3 13 13 14 N N A	Sequence 26, Appl Sequence 27, Appl Sequence 28, Appl Sequence 33, Appl Sequence 571, App Sequence 29, Appl Sequence 28, Appl Sequence 114, App Sequence 334, App Sequence 360, App Sequence 360, App

ALIGNMENTS

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US-09-852-903C-23

Sequence 23, Application US/09852903C

Publication No. US20030104376A1

GENERAL INFORMATION:

APPLICANT: Diatech Pty. Ltd.

TITLE OF INVENTION: An assay

FILE REFERENCE: 2414918/EJH

CURRENT APPLICATION NUMBER: US/09/852,903C

CURRENT FILING DATE: 2001-05-09

PRIOR APPLICATION NUMBER: US 60/202,771

PRIOR APPLICATION NUMBER: US 60/202,771

PRIOR APPLICATION NUMBER: US 60/202,559

PRIOR PILING DATE: 2000-05-10

NUMBER OF SEQ ID NOS: 38

SOFTWARE: Patentin version 3.0

SEQ ID NO 23

LENGTH: 36

TYPE: DNA

ORGANISM: artificial sequence

PEATURE:

NAME/KEY: misc feature

LOCATION: ()...()

OTHER INFORMATION: CA-19

US-09-852-903C-23
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US-09-852-903C-23
                                                                                 Matches
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Best Local
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                        h 100.0%; Score 36; DB 10; Similarity 100.0%; Pred. No. 6.5e-05; 36; Conservative 0; Mismatches 0;
CACACACACACACACACACACACACACACACACACA
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US-09-263-959-678/c
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TITLE OF INVENTION: An assay
FILE REFERENCE: 24141918/EJH
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR TILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR FILING DATE: 2000-05-10
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APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test for
FILE REFERENCE: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                       US-09-852-903C-24
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               Sequence 678, Application US/09263959 Patent No. US20020150891A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
SEQ ID NO 24
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Best Local :
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                                                                                                                                                                                                                                                                                                                   LOCATION: ()..()
OTHER INFORMATION: CA-20
                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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APPLICANT:
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Hood, Leroy E
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100.0%; Pred. No. 6.5e-05;
tive 0; Mismatches 0;
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Pred. No. 6.5e-05;
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CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
SEQ ID NO 25
LENGTH: 40
TYPE: DNA
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                                                                    ; OTHER INFORMATION: CA-21 US-09-852-903C-25
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                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25, Application US/09852903C Publication No. US20030104376A1 GENERAL INFORMATION:
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APPLICATION UMMBER: US/09/262 ^-
FILING DATE: 05-Map.
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Best Local :
Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 36; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Diatech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 2414918/EJH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (206) 682-60
INFORMATION FOR SEQ ID NO:
                                                                                                      ORGANISM: artificial sequence FEATURE:
NAME/KEY: misc feature
LOCATION: ()...()
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: MCMBBECTS, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 92
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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APPLICANT: Koop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH
NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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TELEPHONE: (206)
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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Similarity 100.0%;
36; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (206) 622-4900
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   Score 36; DB 10;
Pred. No. 6.5e-05;
); Mismatches 0;
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Pred. No. 6.5e-05;
); Mismatches 0;
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US-10-661-088-24
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CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-12-05
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
                                                                        US-10-661-097-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24, Application US/10661097 Publication No. US20040162254A1 GENERAL INFORMATION:
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SEQ ID NO 24
LENGTH: 40
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                APPLICANT: JUTEAU, JEAN-MARC
TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING HSV
FILE REFERENCE: 029849/0204
CURRENT APPLICATION NUMBER: US/10/661,097
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR FILING DATE: 2002-09-13
PRIOR FILING DATE: 2002-09-13
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TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING HSV
FILE REFERENCE: 029849/0206
                                                                                                                                                                                                                                                            SOFTWARE: Patentin Ver.
                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: VAILLANT, ANDREW APPLICANT: JUTEAU, JEAN-MAJ
                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: oligonucleotide
                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                              ENGTH: 40
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Similarity 100.0%; Pred. No. 6.5e-05;
36; Conservative 0; Mismatches 0;
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Score 36;
Pred. No.
DB 18;
6.5e-05;
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                     Length 40
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                                      US-10-661-099-24
                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/661,099
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver: 3.2
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US-10-661-355-24
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CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
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Publication No. US20040171568A1
GENERAL INFORMATION:
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Publication No. US20040170959A1
GENERAL INFORMATION:
APPLICANT: VAILLANT, ANDREW
APPLICANT: JUTEAU, JEAN-MARC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 24
Query Match
                                                                                                                                                                          SEQ ID NO 24
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Best Local (
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TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES
FILE REFERENCE: 029849/0208
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: JUTEAU, JEAN-MARC
TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING HIV
                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 029849/0203
                                                          OTHER INFORMATION:
OTHER INFORMATION:
                                                                                               LENGTH: 40
TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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                                                      Description of Artificial Sequence: Synthetic oligonucleotide
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  100.0%; Score 36;
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6.5e-05;
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  18;
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Length 40;
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US-10-661-415-24
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CURRENT APPLICATION NUMBER: US/10/661,415

CURRENT FILING DATE: 2003-09-12

PRIOR APPLICATION NUMBER: PCT/IB03/04573

PRIOR FILING DATE: 2003-09-11

PRIOR APPLICATION NUMBER: 60/430,934

PRIOR FILING DATE: 2002-12-05

PRIOR PFILICATION NUMBER: 60/410,264

PRIOR APPLICATION NUMBER: 60/410,264

PRIOR APPLICATION NUMBER: 60/410,264

PRIOR FILING DATE: 2002-09-13

NUMBER OF SEQ ID NOS: 36
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Best Local Similarity
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Publication No. US20040229828A1
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TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING RSV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: VAILLANT, ANDREW APPLICANT: JUTEAU, JEAN-MAI
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                                                     ZIF: 98104-7092
ZIF: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
FILING DATE: 05-MAR-1999
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Hood, Leroy E.

APPLICANT: Rowen, Lee

APPLICANT: KOOD, Ben F.

TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI

NUMBER OF SEQUENCES: 1279
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                                                                                                                                                                                                                                                                                     STREET: 6300 (
CITY: Seattle
                                        CLASSIFICATION:
                                                                                                                                                                                                                                              COUNTRY: US
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                                                                                                                                                                                                                                                                 Washington
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6300 Columbia Center, 701 Fifth Avenue
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Sequence 27, Application US/09852903C
Publication No. US20030104376A1
GENERAL INFORMATION:
APPLICANT: Diatech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 2414918/EM.
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
PRIOR FILING DATE: 2000-05-10
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US-09-852-903C-27
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US-09-852-903C-26
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Best Local (
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SEQ ID NO 26
LENGTH: 42
TYPE: DNA
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CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09
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PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
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SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: CA-22
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NAME/KEY: misc feature
LOCATION: ()..()
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
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TYPE: nucleic acid
STRANDEDNESS: single
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Pred. No. 6.5e-05;
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Sequence 28, Application US/09852903C
Publication No. US20030104376A1
GENERAL INFORMATION:
APPLICANT: Diatech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 2414918/EJH
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR PILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
SEQ ID NO 28
LENGTH: 46
TYPE: DNA
ROANISM: artificial sequence
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, NAME/KEY: misc feature
; LOCATION: ()..\(7\)
; OTHER INFORMATION: CA-23
US-09-852-903C-27
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US-09-971-353-33
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                                                              Sequence 33, Application US/09971353
Publication No. US20030113723A1
GENERAL INFORMATION:
APPLICANT: Bapat, Bharati
APPLICANT: Rose, Melanie Anne
TITLE OF INVENTION: METHOD FOR EVALUATING MICROSATELLITE INSTABILITY IN A TUMOR SAMPL
FILE REFERENCE: 11757.54USU1
CURRENT FILING DATE: 2001-10-04
PRIOR APPLICATION NUMBER: US 60/237,884
PRIOR APPLICATION NUMBER: US 60/237,884
PRIOR FILING DATE: 2000-10-04
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PATENTIN VETSION 3.1
SEQ ID NO 33
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Best Local S
Matches 36
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LENGTH: 44
TYPE: DNA
ORGANISM: artificial sequence
LENGTH: 46
TYPE: DNA
ORGANISM: Homo mapiens
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NAME/KEY: misc_feature
LOCATION: ()..()
OTHER INFORMATION: CA-24
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